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<b>(21) International Application Number:</b> PCT/GB97/01161  <b>(22) International Filing Date:</b> 25 April 1997 (25.04.97)  <b>(30) Priority Data:</b> 9608626.9                      25 April 1996 (25.04.96)                      GB  <b>(71) Applicant (for all designated States except US):</b> MUREX DIAGNOSTICS CORPORATION [BB/BB]; Whitepark House, White Park Road, Bridgetown, Barbados (BB).  <b>(72) Inventors; and</b> <b>(75) Inventors/Applicants (for US only):</b> TEDDER, Richard, Seton [GB/GB]; UCL Medical School, Windeyer Building, Dept. of Virology, 46 Cleveland Street, London W1P 6DB (GB). IJAZ, Samreen [GB/GB]; UCL Medical School, Windeyer Building, Dept. of Virology, 46 Cleveland Street, London W1P 6DB (GB). FERNS, Ruth, Bridget [GB/GB]; UCL Medical School, Windeyer Building, Dept. of Virology, 46 Cleveland Street, London W1P 6DB (GB).  <b>(74) Agents:</b> SILVESTON, Judith et al.; Abel & Imray, Northum- berland House, 303-306 High Holborn, London WC1V 7LH (GB).	<b>(81) Designated States:</b> AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, DE, DK, EE, ES, FI, GB, GE, GH, HU, IL, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, TJ, TM, TR, TT, UA, UG, US, UZ, VN, YU, ARIPO patent (GH, KE, LS, MW, SD, SZ, UG), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, ML, MR, NE, SN, TD, TG).  <b>Published</b> <i>With international search report.</i> <i>Before the expiration of the time limit for amending the claims and to be republished in the event of the receipt of amendments.</i>	
<b>(54) Title:</b> HEPATITIS B MONOCLONAL ANTIBODIES  <b>(57) Abstract</b>  A monoclonal antibody that is capable of binding specifically to wild type HBsAg and to at least two mutant forms of HBsAg may be used in an improved immunoassay for the detection of both escape mutants and wild type HBsAg and may be used for passive immunisation against HBV.		

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HEPATITIS B MONOCLONAL ANTIBODIES

On a global scale, the hepatitis B virus (HBV) is the most significant of the hepatotropic viruses in terms of the number of people chronically infected and the severity of the complications of infection.

Hepatitis B emerged as a major problem after the introduction of parenteral therapy, mass immunisation campaigns and the extensive transfusion of blood and pooled blood products. Long incubation periods, high incidence of asymptomatic infections and the occurrences of an infectious carrier state make HBV well suited to transmission by blood. Numerous workers have also provided evidence for sexual transmission of HBV, and transmission of hepatitis B virus from carrier mothers to their babies can occur during the perinatal period.

HBV, a DNA virus, is a member of the hepadna virus group. Electron microscopy of the HBV reveals a double shelled, spherical particle of 42nm in diameter called the "Dane particle". Those virus particles exhibit an electron dense spherical inner core with a diameter of approximately 22-25nm and an outer coat 7nm in thickness. The outer coat of the virus bears the surface antigen (called herein "HBs" or "HBsAg") to which virus-neutralising antibody is directed. The spherical 20nm diameter inner-core particles bear the viral core antigen (HBcAg) and the e antigen (HBeAg), the viral DNA, DNA polymerase activity and a protein kinase activity.

HBsAg contains the major neutralising epitope of HBV, termed the "a" determinant, which spans amino acids 124 to 147 and is common to all HBV isolates, see for example, Pugh et al (1986). The development of anti-HBs after acute or chronic HBV infection is usually associated with recovery and a good prognosis. Anti-HBs is also associated with the production of neutralising antibodies resulting from vaccination. The majority of anti-HBs found in convalescent and post-vaccination sera binds in the region of the "a" determinant, which has as yet undefined structure. It is clear that the "a" determinant is highly conformational because

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denaturation of that area by alkylation or reduction gives rise to HBsAg particles with greatly reduced antigenicity. It is thought that disulphide bridges between cysteine residues are responsible for correct conformation. One possible  
5 structure of the "a" determinant involves a disulphide bridge between amino acids 124 and 137, forming a first loop, and a further disulphide bridge between amino acids 139 and 147, forming a second loop.

Anti-HBs is thought to bind predominantly to the second  
10 loop but it is thought that the epitopes are not confined to one loop only. The whole of the sequence of the "a" determinant probably contributes to the antigenic structure. The "a" determinant is conserved though there is a degree of amino acid variation in normal isolates of HBV. Greater  
15 variation is accepted in the first loop of the putative epitope, perhaps implying that this region does not contribute significantly to the neutralisation epitope.

The "a" determinant is found in all subtypes of HBV and it is variation of amino acids within and around the  
20 "a" determinant that gives rise to subtypes. HBsAg can be classified into four major immunological subtypes, adw, ayw, adr, and ayr, each with an associated geographical distribution. The d/y and w/r subtypes are determined by substitutions of lysine by arginine at amino acids 122 and  
25 160 respectively.

Recently, the presence of HBsAg has been observed in anti-HBs serum samples of certain patients. The HBsAg in those patients may not be neutralised by the anti-HBs present, implying the presence of HBsAg variants.  
30 Significant variants (mutants) have been associated with vaccination, monoclonal antibody therapy, polyclonal antibody therapy and cases of HBV infection difficult to diagnose in clinical laboratories see, for example, Carmen et al. (1992, 1993), Harrison et al. (1993), Hawkins et al. (1994), Howard  
35 et al. (1993), McMahon et al. (1992), Okamoto et al. (1992), and Wallace et al. (1994).

Upon analysis the mutants seem to have been selected

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from a mixed population and to have point mutations causing amino acid substitutions in the "a" determinant. The mutants are thought to arise by random mutations within the gene which gives rise to a pool of genotypes. In the mutants 5 described to date, it is thought that the immune response is the predominant factor in selecting for the mutants.

Generally, addition of monoclonal antibodies to virus infected cells in vitro result in selection of isolates that are not neutralised by that antibody. It is thus not 10 surprising that monoclonal antibodies given to patients with active viral replication can result in selection of so-called "escape" mutants.

Several separate escape mutants with clinical significance have been described in vaccinated individuals. 15 In three cases there was found to be a point mutation at the codon for amino acid 145 in the "a" determinant of HBsAg resulting in an amino acid change from glycine to arginine. Administration of serum containing one such mutant virus to a chimpanzee proved that the agents are transmissible.

20 In a mutant virus implicated in a breakthrough infection in a vaccinated population, a lysine to glutamic acid mutation at amino acid 141 of the "a" determinant was found. That and other point mutations resulting in one or more amino acid substitutions in the "a" determinant have 25 since been reported in vaccinated individuals.

Escape mutants are a cause for concern on several accounts. Firstly, there is failure to detect such mutants by immunoassays. Diagnostic assays are designed to achieve high sensitivity and specificity. Assays for HBsAg 30 detection depend on interaction between an anti-HBs reagent and HBsAg in the sample under investigation. A resulting anti-HBs/HBsAg complex is then detected. If there is a significant mutation in the HBsAg epitope and it is not recognised by the anti-HBs, then the HBsAg will either not be 35 detected or the assay will be very insensitive. Failure to detect an escape mutant may not only affect the person harbouring the mutant; it may lead to transmission of

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infection through donated blood, blood products or organs. Secondly, HBV with mutant HBsAg may infect individuals even though they have been previously immunised and have an anti-HBs response.

5 A monoclonal antibody to an escape mutant at position 122 has been described in WO94/26904. That antibody is able to discriminate between the wild type HBsAg and the mutant form and hence enables identification of the mutant form.

The present invention provides a monoclonal antibody  
10 that is capable of binding specifically to wild-type HBsAg and to at least two, preferably more than two, mutant forms of HBsAg.

The ability of a monoclonal antibody of the present invention to bind specifically to both wild type and mutant  
15 HBsAg suggests that it is binding to a region of the surface antigen that is conserved between the wild type protein and the mutant forms. The conserved region and hence the antigenic determinant (or epitope) may be in the "a" determinant itself, in the region of the "a" determinant or  
20 even in a non-"a" region.

Because of its ability to bind specifically to mutant forms of HBsAg as well as to the wild type protein, and hence to detect escape mutants, a monoclonal antibody of the present invention is particularly useful in improving the  
25 efficacy of immunoassays for the detection of HBsAg, for clinical diagnosis of HBV infections and also for blood screening. The safety of the blood supply and of the supply of blood products may be improved by the use of a monoclonal antibody of the invention in an HBsAg assay, either alone,  
30 that is to say, as the only anti-HBs antibody or, especially, in addition to one or more other anti-HBs antibodies.

Figures 1a and 1b of the accompanying drawings set out amino acid and nucleic acid sequences of part of the HBV surface region encompassing the "a" determinant. Figure 1b  
35 of the drawings as filed is a typescript version of Figure 1a. In each of Figures 1a and 1b:  
Line 1 sets out recognised amino acid variants of subtype

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adyw; Line 2 sets out the consensus amino acid sequence of subtype adyw; Line 3 sets out the consensus nucleic acid sequence of subtype adyw; Line 4 sets out recognised nucleic acids variants; Line 5 sets out the nucleic acid variants encoding Mutant HBsAg II (MAM HBsAg); and Line 6 sets out the nucleic acid variants encoding Mutant HBsAg I (NP HBsAg).

A further recognised nucleotide variant at position 143 is ACG for TCG, which results in threonine instead of serine at that position.

10 A putative monoclonal antibody of the present invention may be screened for the ability to bind specifically to three or more reference antigens namely wild type HBsAg and two or more mutant forms of HBsAg. A skilled worker is able readily to distinguish specific binding between an antibody and an  
15 antigen from non-specific binding. Any antibody that is capable of binding specifically to three or more such reference antigens is a monoclonal antibody of the present invention. Such a method of screening is itself part of the present invention. It should be noted that the term  
20 "binding" is used throughout the present specification to denote specific binding.

An HBsAg used for screening and/or as an antigen in the production of hybridomas and monoclonal antibodies of the invention may be a full length protein or may be an  
25 appropriate antigenic fragment or derivative of a wild type or mutant HBsAg.

A monoclonal antibody of the present invention may be of any immunoglobulin class, for example, IgG, IgM or IgA, and of any isotype.

30 Mutant forms of HBsAg may be found in mutant viruses having a mutation that leads to at least one amino acid substitution relative to wild type HBsAg, for example, as found in so-called "escape mutants". A mutant HBsAg may have a substitution in the "a" determinant or in the region  
35 of the "a" determinant, for example, the mutation may be a point mutation. A mutation, for example, a point mutation, may be within the sequence encoding amino acids 133 to 145 of

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HBsAg. The mutation may lead to an amino acid substitution at position 133 and/or at position 145. Further and/or different substitutions may be present, for example, at any one or more of positions 134, 141, 142, 143 and 144. A mutant  
5 HBsAg may, for example, have amino acid substitutions relative to the wild type at any one or more of positions 143, 144 and 145. Such mutations may be present in addition to other mutations, either in the "a" determinant or in another region.

- 10 A monoclonal antibody of the present invention is capable of binding specifically, for example, to wild-type HBsAg and to two or more different mutant HBsAg as described above, for example, each having a mutation at any one or more of positions 133, 134, 141, 142, 143, 144 and 145.
- 15 Alternatively, binding may be to one mutant HBsAg described above and to a mutant HBsAg having a mutation in a different region.

Examples of mutant forms of HBsAg are those having the following substitutions relative to the wild type in the  
20 region of the "a" determinant:

Mutant HBsAg I ("NP" HBsAg): met to ile at amino acid 133; phe to his at amino acid 134; and asp to val at amino acid 144;

Mutant HBsAg II ("MAM" HBsAg): met to ile at amino acid 133;  
25 phe to asn at amino acid 134; pro to ser at amino acid 142; ser to leu at amino acid 143; and gly to lys at amino acid 145;

Mutant HBsAg III ("SZ" HBsAg): gly to arg at amino acid 145;  
Mutant HBsAg IV ("SP" HBsAg): ser to met at amino acid 143.

- 30 A putative antibody may be screened, for example, against wild type HBsAg and against any two or more of the mutant forms described above. For example, a putative monoclonal antibody of the present invention may be screened against wild-type HBsAg and against two or more different  
35 mutant HBsAg each having mutation(s) at any one or more of positions 133, 134, 141, 142, 143, 144 and 145, especially 143, 144 and 145. Alternatively, one such mutant may be used



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with a mutant HBsAg having a mutation in a different region. Mutant forms I and IV described above may be used for screening.

The present invention provides a monoclonal antibody 5 that is capable of binding specifically to wild-type HBsAg and to at least one mutant HBsAg carrying an "a" determinant coded for by a sequence having point mutations at any one or more of the codons encoding amino acids 143, 144 and 145. The monoclonal antibody may bind to two or more such mutant 10 HBsAg or to one such mutant HBsAg and to another different mutant HBsAg.

Examples of monoclonal antibodies of the present invention are those produced by the hybridomas designated P2D3, M3A10 and M4F5, which hybridomas have been deposited at 15 the European Collection of Cell Structures (ECACC), Virus Research and Production Laboratory, Public Health Laboratory Service, Centre for Applied Microbiology & Research, Porton Down, Salisbury, Wiltshire SP4 0JG, England, under the terms of the Budapest Treaty on the International Recognition of 20 the Deposit of Microorganisms for the Purposes of Patent Procedure, 1977. The accession numbers and dates of the hybridoma deposits are as follows: Hybridoma P2D3: Accession Number ECACC 97042331, Accession Date 23 April 1997; Hybridoma M3A10: Accession Number ECACC 97042330, Accession 25 Date 23 April 1997; Hybridoma M4F5: Accession Number ECACC 97042519, Accession Date 25 April 1997.

In this specification, a monoclonal antibody may be referred to under the same designation as the hybridoma that produces it, for example, the monoclonal antibody designated 30 P2D3 is as produced by hybridoma P2D3 (ECACC 97042331). Hybridomas P2D3 (ECACC 97042331), M3A10 (ECACC 97042330) and M4F5 (ECACC 97042519) and the monoclonal antibodies they produce are described in detail in the Examples below.

Each of the monoclonal antibodies P2D3, M3A10 and M4F5 35 is capable of binding specifically to (i) wild type HBsAg, (ii) Mutant HBsAg I (NP HBsAg) and (iii) Mutant HBsAg II (MAM HBsAg). Furthermore, each also is capable of binding

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specifically to Mutant HBsAg III (SZ HBsAg) and to Mutant HBsAg IV (SP HBsAg). Monoclonal antibodies P2D3 and M4F5 are IgG antibodies. Monoclonal antibody M3A10 is an IgA antibody.

5 It is known that wild type HBsAg has a number of different serotypes, and also that there are recognised mutants see, for example, Pugh et al 1980. The term "wild type HBsAg" as used in the present specification includes all HBsAg that are recognised in the art as wild type or that  
10 would be acknowledged to be wild type. The term therefore includes wild type HBsAg of all serotypes and of all recognised variants. The term "mutant HBsAg" as used herein does not include recognised wild type variants.

Suitable screening tests are well known to those  
15 skilled in the art and include cross-competition assays and assays, for example, using radio-labelled monoclonal antibody to determine binding specificities.

As stated above, the ability of a monoclonal antibody of the present invention to bind to both wild type and mutant  
20 HBsAg suggests that it is binding to a region of the surface antigen that is conserved between the wild type protein and the mutant forms. (See also the General Discussion below.) The location and the sequence of such a conserved region and hence a conserved epitope may be determined by known  
25 methods, for example, by epitope mapping optionally using a monoclonal antibody of the invention. Epitopic peptides and polypeptides may then be produced, for example, recombinantly, by chemical synthesis or by a combination of different methods. A resulting antigenic peptide or  
30 polypeptide may also be immunogenic. An epitope against which a monoclonal antibody of the invention is directed is itself part of the present invention.

The present invention includes fragments and derivatives of a monoclonal antibody of the present invention,  
35 for example, Fab and Fab<sub>2</sub> fragments and conjugates. Derivatives include humanised derivatives. Methods for producing fragments and derivatives, including humanised

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derivatives, are well known. For example, fragments and derivatives may be produced recombinantly. Fragments and derivatives of antibodies and their uses are well known to the skilled worker. The term "monoclonal antibody" as used 5 herein includes fragments and derivatives thereof.

The present invention also provides an anti-idiotypic antibody to a monoclonal antibody of the present invention. Such antibodies, which comprises an "internal image" of the original epitope, may be used as epitope substitutes and are 10 particularly useful in the case of conformational epitopes, since it is difficult to map such epitopes and it may not be possible to produce synthetic epitopic peptides in such cases.

A monoclonal antibody of the present invention may be 15 used in an immunoassay for the detection of HBsAg ("HBV assay") and hence of hepatitis B infections, either as a substitute for or, especially, in addition to anti-HBs antibodies currently used in HBV assays. Such assays may be used for clinical diagnosis or for blood screening, and the 20 present invention includes both methods of diagnosis and methods of blood screening using an immunoassay comprising a monoclonal antibody of the present invention.

A monoclonal antibody of the present invention may also be used therapeutically or prophylactically as an antiserum 25 for passive immunisation and/or to define an epitope for use as a vaccine in active immunisation. The present invention includes such antisera and methods of immunisation, and also epitopes defined using such antisera.

A further application of a monoclonal antibody of the 30 present invention is in affinity chromatography, for example, to purify wild or mutant types of HBsAg, antigenic fragments thereof, antigenic peptides or anti-idiotypic antibodies.

Accordingly, the present invention provides an immunoassay for the detection of HBsAg, which comprises 35 contacting a sample under investigation with a monoclonal antibody of the present invention, a fragment or derivative thereof, or a combination of two or more thereof, and

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detecting any resulting antigen-antibody complex.

The term "detection" is used herein to denote detection and/or determination and includes qualitative, quantitative and semi-quantitative methods.

5       The monoclonal antibody of the invention, fragment, derivative or combination thereof may be used with one or more other antibodies selected from polyclonal anti-HBs antibodies and other monoclonal anti-HBs antibodies.

10       An immunoassay of the present invention may be in a homogeneous or heterogeneous format. The format may be a capture or a competitive format.

15       An immunoassay for the detection of antibodies to hepatitis B core antigen (anti-HBc) may be carried out simultaneously with the assay for HBsAg. HBV combination anti-core/surface antigen assays are known and are in use.

20       The present invention further provides an immunoassay kit that comprises a monoclonal antibody of the present invention or a fragment or derivative thereof, or a combination of two or more thereof, and other reagents required for carrying out an immunoassay for HBsAg and optionally also reagents for detecting anti-HBc antibodies. One or more other anti-HBs antibodies selected from polyclonal anti-HBs antibodies and other monoclonal anti-HBs antibodies may be present, and the other reagents may be  
25       selected from washing solutions, diluents, standard solutions, control reagents and labelled anti-HBs antibodies.

30       The present invention further provides a solid phase suitable for use in an immunoassay on which is immobilised a monoclonal antibody of the present invention, a fragment or derivative thereof, or a combination of two or more thereof. One or more further anti-HBs antibodies selected from polyclonal anti-HBs antibodies and other monoclonal anti-HBs antibodies may also immobilised on the solid phase. Furthermore, an agent capable of capturing anti-HBc  
35       antibodies may be immobilised on the solid phase in addition to the anti-HBs antibodies.

      The present invention further provides an antiserum

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suitable for use therapeutically or prophylactically for passive immunisation which comprises a monoclonal antibody of the present invention, a fragment or derivative thereof, or a combination of two or more thereof.

5       The present invention also provides a composition suitable for use therapeutically or prophylactically for passive immunisation against HBV which comprises a monoclonal antibody of the present invention, a fragment or derivative thereof, or a combination of two or more thereof, in  
10 admixture with a pharmaceutically suitable carrier.

An antiserum or composition of the invention may also comprises one or more other antibodies selected from polyclonal anti-HBs antibodies and other monoclonal anti-HBs antibodies.

15       The present invention further provides a method of therapeutic or prophylactic passive immunisation against HBV which comprises administering to a human a therapeutically or prophylactically effective amount of a monoclonal antibody of the invention, a fragment or derivative thereof or a  
20 combination of two or more thereof. One or more other antibodies selected from monoclonal and polyclonal anti-HBs antibodies may also be administered.

The present invention further provides isolated Mutant HBsAG I, Mutant HBsAG II, Mutant HBsAG III or Mutant HBsAG IV  
25 as defined above. The invention also provides a fragment or derivative of such a mutant HBsAg, said fragment or derivative being capable of binding specifically to a monoclonal antibody of the present invention or to a fragment or derivative thereof. A fragment or derivative of a mutant  
30 HBsAg may also be immunogenic. A mutant HBsAg or a fragment or derivative thereof is obtainable from a natural source or recombinantly. It may be preferable, for example, for use as an immunogen, to have the mutant HBsAg or fragment or derivative thereof in partially or fully glycosylated form.  
35 The present invention also provides a chemically synthesised antigenic and optionally also immunogenic epitopic peptide.

The present invention provides an vaccine

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composition that comprises a mutant HBsAg of the present invention or an immunogenic fragment or derivative thereof, or a combination of two or more thereof, and a pharmaceutically suitable carrier. A vaccine composition may  
5 further comprise one or more other immunogenic peptides or polypeptides, for example, other immunogenic HBsAg peptides or polypeptides.

The present invention also provides a method of vaccinating a human against HBV, which comprises  
10 administering to the human a therapeutically effective amount of a mutant HBsAg of the present invention, an immunogenic fragment or derivative thereof, or a combination of two or more thereof, optionally in combination with one or more other immunogenic HBsAg peptides or polypeptides.

15 The present invention also provides an isolated nucleic acid that encodes Mutant HBsAG I, Mutant HBsAG II, Mutant HBsAG III or Mutant HBsAG IV as defined above, or a fragment or derivative of such a mutant HBsAg, said fragment or derivative being capable of binding specifically to a  
20 monoclonal antibody of the present invention or to a fragment or derivative thereof. The fragment or derivative of the mutant HBsAg is preferably immunogenic.

The present invention also provides a composition that comprises a nucleic acid of the invention, especially a  
25 vaccine composition that comprises a nucleic acid of the invention in a form suitable for expression of the encoded amino acid sequence when administered to a mammal.

For any of the applications and uses according to the present invention, the monoclonal antibody of the present  
30 invention may be monoclonal antibody P2D3, monoclonal antibody M3A10, monoclonal antibody M4F5 or a combination of two or more thereof. Any one or more of the monoclonal antibodies P2D3, M3A10 and M4F5 may be used together with any other monoclonal antibody of the present invention and/or  
35 with any other anti-HBV antibodies, especially antiHBS antibodies. A fragment or derivative of a monoclonal antibody of the invention may be used.

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The present invention also provides a hybridoma capable of producing a monoclonal antibody of the present invention, for example, hybridoma P2D3 (Accession Number ECACC 97042331), hybridoma M3A10 (Accession Number ECACC 97042330) 5 and hybridoma M4F5 (Accession Number ECACC 97042519).

The present invention provides a method of producing a hybridoma capable of producing a monoclonal antibody of the present invention, which comprises immunising an animal with wild type HBsAg or a mutant form of HBsAg, immortalising 10 antibody producing cells to form hybridomas, screening the resulting hybridoma culture against wild type HBsAg and two or more mutant forms of HBsAg, and selecting those hybridomas that produce a monoclonal antibody that is capable of binding specifically to wild type HBsAg and two or more mutant forms 15 of HBsAg.

A monoclonal antibody of the invention may be obtained by culturing a hybridoma of the invention in vitro or in vivo and obtaining the monoclonal antibody from the culture medium. In the case of in vivo production, the culture 20 medium is ascitic fluid.

Methods for raising and for screening hybridomas and monoclonal antibodies are well known to those skilled in the art. The general methodology is set out by Kohler & Milstein (1975) and there have been numerous publications on the 25 subject since then. Any such method may be used to raise and to screen hybridomas and monoclonal antibodies according to a method of the present invention. The antigen used to raise the antibodies may be wild type HBsAg or a mutant form of HBsAg, for example, as described above, for example, one or 30 more of the mutant forms I to IV described above. The screening of the resulting hybridomas and monoclonal antibodies for their ability to bind specifically to different forms of HBsAg may be carried out by any conventional method using wild type HBsAg and the chosen 35 mutant forms. As described above, screening may be carried out using, as reference antigens, wild type HBsAg and two or more mutant forms of HBsAg, for example, as described above, for

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example, two or more of the mutant forms I to IV.

For screening, it is particularly convenient to use an IgG reverse capture radioimmunoassay, which is capable of a very low background and a large dynamic range, and has the further advantage of being simple to set up.

In an IgG reverse capture assay for an antibody of interest, an appropriate anti-species IgG is immobilised on a solid phase, for example, microwells. If, for example, the antibodies under investigation have been raised in mice, then the solid phase is coated with anti-mouse IgG. The sample under investigation is then incubated with the immobilised anti-species IgG, whereupon a representative proportion the total IgGs present in the sample are captured. The solid phase is then incubated with an appropriate labelled antigen, which forms a complex only with the captured antibody of interest. Any complexes formed may then be detected.

In the present case, a sample under investigation, for example, a tissue culture supernatant fluid from a hybridoma culture or ascitic fluid, is incubated with immobilised anti-IgG. The captured antibodies are incubated separately with labelled wild type HBsAg and with two or more labelled mutant HBsAg, for example, as described above, for example, selected from mutant forms I to IV. Any sample that is capable of binding specifically wild type HBsAg and two or more mutant surface antigens contains a monoclonal antibody that falls within the present invention.

A reverse phase IgM or IgA assay may be used analogously to obtain an IgM or IgA monoclonal antibody.

Commercially available immunoassays for HBsAg are generally heterogeneous phase or "capture" assays, in which anti-HBs is immobilised on a solid phase, usually microwells, small particles or beads. When a sample under investigation, generally serum, is contacted with the immobilised anti-HBs, HBsAg present in the sample should bind to the immobilised antibody. The captured antigen is then detected, generally using a labelled anti-HBs. In conventional assays, the antibody used for capture or for



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detection may be monoclonal or polyclonal.

Polyclonal antibodies, because of their nature, have a range of specificity and selectivity. In the case of HBV, this may result in the presence in the antiserum of an antibody that can bind to an escape mutant. However, that cannot be guaranteed, and there is an inherent problem in obtaining uniformity of product from batch to batch. Monoclonal antibodies have defined specificity and selectivity and may therefore fail to detect escape mutants.

10 The use of a monoclonal antibody of the present invention, which is capable of binding specifically to both wild type HBsAg and to mutant HBsAg, or a combination of two or more such antibodies, including fragments and derivatives thereof, in an immunoassay for HBsAg improves the performance of the assay and reduces the possibility that mutant HBsAg will not be detected. A monoclonal antibody of the present invention or a combination of two or more such antibodies may be used either as the only anti-HBs antibody reagent or, preferably, may be used in addition to polyclonal anti-HBs or other monoclonal anti-HBs, for antigen capture and/or for detection of any resulting antibody-antigen complex. A monoclonal antibody of the present invention, or a combination of two or more thereof, may be used analogously to conventional anti-HBs in a homogeneous phase assay for HBsAg.

25 The present invention accordingly provides an immunoassay for the detection of HBsAg, which comprises contacting a sample suspected of containing HBsAg with a monoclonal antibody of the present invention, a fragment or derivative thereof, or a combination of two or more thereof, and detecting any resulting antigen-antibody complex. The monoclonal antibody, fragment, derivative or combination thereof may be used as the only anti-HBs reagent or may be used with one or more other anti-HBs antibodies selected from polyclonal anti-HBs and other anti-HBs monoclonal antibodies.

35 Immunoassays of various different formats, methods for carrying them out and suitable reagents are a well known and are described in various textbooks and review articles, for

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example, Kemeny & Challacome 1988 and Tsu & Herzenberg 1980. Any method for the detection of an antigen may be used in the present invention.

An assay according to the present invention may be a  
5 so-called "sandwich" assay, a competition assay or a direct reaction. The monoclonal antibody, fragment, derivative or combination thereof may be immobilised on a solid surface alone or in admixture with a polyclonal anti-HBs antiserum and/or with one or more other anti-HBs monoclonal antibodies.  
10 Alternatively the monoclonal antibody of the invention or the combination thereof and any further antibodies may be in a homogeneous phase. Any resulting antibody-antigen complex may be detected using labelled anti-HBs. In the case of a heterogeneous phase assay, the anti-HBs used for detection  
15 may be the same as or different from that used for coating the solid surface.

Suitable solid surfaces on which anti-HBs may be immobilized are well known and include the inner walls of wells of microtitre plates, beads, particles and so-called  
20 "latex". Membranes and strips, for example, of nitrocellulose or paper may be used. A membranes or strip may be incorporated in an assay device. Such devices are also well known. The invention includes such solid phases on which a monoclonal antibody of the present invention, or a fragment  
25 or derivative thereof, or a combination of two or more thereof is immobilised.

The anti-HBs used for detection of any antigen-antibody complex may be labelled with any agent capable of generating a signal directly or indirectly. Such agents are well known.  
30 Agents capable of generating a direct signal include radiolabels, chromogenic and fluorescent labels. Agents capable of generating a signal indirectly include enzymes capable of catalysing a reaction that gives rise to a colour change.

35 The present invention also provides a kit which comprises the components requires to carry out an immunoassay of the present invention. Such a kit comprises, for example,

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a monoclonal antibody of the present invention or a combination of two or more thereof immobilised on a solid surface, together with container(s) comprising other reagent(s) required, for example, selected from washing solutions and diluents, standard solutions and control reagents, labelled anti-HBs antibodies and, in the case of enzyme-labelled anti-HBs, colour reagents.

Often an immunoassay for antibodies to hepatitis B core protein (HBc) is carried out simultaneously with the assay for HBs, for example, in the same microwell. The present invention includes the use of a monoclonal antibody of the present invention in such assays, and kits and coated solid phases for use with such assays.

As indicated above, a monoclonal antibody of the present invention may be used for passive immunisation, for example, during liver transplantation of an HBV infected patient. For passive immunisation, the antibody is brought into a suitable form for use as an antiserum for parenteral administration, for example, in a form allowing derivatisation including humanisation. The dose of antibody to be used will depend on the particular circumstances of each patient and will be determined on a case-by-case basis.

A monoclonal antibody of the present invention, including combinations of two more such antibodies, fragments and derivatives thereof, may be used alone or in addition to other anti-HBs antibodies, either polyclonal or monoclonal. Other anti-HBV antibodies may also be included.

A mutant form of HBsAg to which a monoclonal antibody of the present invention binds may itself be used as an antigen, for example, as a vaccine; in the production of monoclonal antibodies; and for screening for putative antibodies according to the present invention. Such a mutant HBsAg may be obtained from a subject believed to be harbouring an HBV escape mutant. Such a subject will generally be both anti-HBs positive and HBsAg positive. Suitable subjects may be found among patients who have received monoclonal antibody therapy for HBV infection, in

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vaccinated subjects, especially where breakthrough infection is found, and in cases of HBV infection that are difficult to diagnose in clinical laboratories using current standard tests.

- 5       A mutant form of HBsAg, for example, one of the mutant forms described above, for example, mutant forms I to IV, may be produced by recombinant DNA technology. Naturally occurring DNA encoding the mutant form may be inserted into an appropriate host for expression of the mutant HBsAg.
- 10- Alternatively, wild type DNA may be modified, for example, by site-directed mutagenesis, and then used to obtain expression of a mutant form.

HBsAg obtained from natural sources, generally from the blood of a selected subject, will have natural glycosylation, and it is generally preferable to use such HBsAg as an antigen. Methods of purifying HBsAg from blood are well known, see for example, Cameron et al. 1980. It may be preferable to produce a recombinant mutant HBsAg that is partially or fully glycosylated.

- 20       The HBsAg may be full-length or an appropriate antigenic fragment may be used. Such a fragment may be obtained by recombinant technology. Alternatively, a fragment that is an antigenic peptide, for example, produced by chemical synthesis may be used. For use as a vaccine, an antigen should be immunogenic ie capable of producing a protective response.

For use in a vaccine, the desired antigen (whether full length, fragment or peptide), and whether obtained recombinantly, synthetically or from natural sources) must be purified and brought into a suitable form, for example, using conventional carriers and, where appropriate, adjuvants.

- 30       As an alternative to using an antigen for vaccination, a nucleic acid encoding the antigen may be used. Methods for presenting a nucleic acid in a form from which it can be expressed in vivo are known. The present invention includes a nucleic acid vaccine.

Methods for producing anti-idiotypic antibodies are well

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known. In the present case a monoclonal antibody of the present invention is used as the antigen in the production of an anti-idiotypic antibody, and the resulting antibodies may be screened against a monoclonal antibody of the present invention. An anti-idiotypic antibody of the present invention may itself be used as a vaccine, in known manner.

The following non-limiting Example illustrates the present invention.

## 10 EXAMPLE

### I. MATERIALS AND METHODS

#### 1: Patients

The monoclonal antibodies described below were raised against the mutant HBsAg of two patients. Those patients were identified as a result of inconsistent serological markers.

Patient MAM was an HBV carrier. The serum was initially, in 1986, HBsAg positive by the polyclonal based reverse passive haemagglutination assay (RPH) but negative by a monoclonal enzyme immunoassay (EIA). In 1988 the patient underwent renal transplantation. When tested again, HBsAg became detectable by both polyclonal and monoclonal antibody based assays but reverted to being detectable by polyclonal antibody assay only. The patient's serum was anti-HBc and HBsAg positive throughout the period of follow-up.

Patient NP underwent renal transplantation in 1985. In 1990 the patient was anti-HBc sero-positive but HBsAg and anti-HBs were not detected. Haemodialysis was started in 1993 and later that year HBsAg was detected by polyclonal based assays only.

Neither of the patients had been previously vaccinated.

The sequences of the HBsAg genes around the "a" determinant were determined by single stranded sequencing carried out directly on PCR products as is described in Hawkins et al., 1994.

#### 2: Production Of Monoclonal Antibodies

##### 2.1: Animals

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Female Balb/c mice used.

## 2.2: Purification of HBsAg

Purification of HBsAg required for immunisation of mice  
5 was based on the method described by Cameron et al., 1980.

High titre HBsAg positive serum was fractionated by  
passing 20ml volumes through a 100x5 cm diameter column of  
Sephadex 6B (Pharmacia Ltd) equilibrated with 0.9% NaCl  
pH 7.6 buffered with Tris and containing 0.1% sodium azide  
10 (Tris/Sal/Az). The HBsAg rich fractions, now free of albumin  
and IgG, were concentrated to a volume of 9ml by  
ultrafiltration in a stirred 200ml ultrafiltration cell  
fitted with an XM 100 A membrane (Amicon Ltd.). To this 2.69g  
of solid CsCl were added and the volume adjusted to 10ml to  
15 give a density of 1.2g/cm<sup>3</sup>. The material was then ultra-  
centrifuged for three days at 124000g in two 5ml tubes in a  
SW50-L swinging bucket rotor (Beckman RIIC Ltd) at 20°C to  
give isopycnic banding in the self forming density gradient.

The main HBsAg band containing the 22nm small particles  
20 was removed from both tubes, pooled and made up to 5ml with a  
solution of CsCl in Tris/Sal/Az (2.885g plus 10ml) which gave  
a final density of 1.2g/cm<sup>3</sup>. The material was ultra-  
centrifuged under the same conditions as before in one 5ml  
tube to give the second isopycnic banding. The final HBsAg  
25 band was removed and dialysed in PBS to remove CsCl.

## 3: Immunisation of animals

Fifty microlitres of purified NP HBsAg mixed in an  
equal volume of Titermax adjuvant (Vaxcel, Inc.) were  
30 injected subcutaneously into a female Balb/c mouse.  
Approximately two months later a dose of 30 µl was given  
intraperitoneally (i.p) followed three months later by a 75µl  
dose also given i.p.. The final 25 µl of HBsAg in saline  
were given intravenously three days before the fusion.

35 The same protocol was used when immunising the mouse  
with the MAM HBsAg.

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#### 4: Culture of Myeloma cells

JK cells derived from P3-X63-Ag8653 (Kearney et al., 1979) were cultured in complete medium (see reagents) at 37°C in a 5% CO<sub>2</sub> in air atmosphere at 100% humidity.

- 5 The cells were split 1:2 approximately three times weekly and maintained at a concentration of between  $2 \times 10^5$  and  $2 \times 10^6$  cells per ml. Only cells in the log phase of growth were used for fusion to splenocytes.

For the fusion, the myeloma cells were spun at 1,500g for 10 minutes and washed three times in the same volume of incomplete medium (see Reagents). The resuspended JK cells were diluted 1:10 in incomplete medium and the viability and concentration of the cells determined by counting the cells using a Neubauer chamber. The concentration of JK cells was 15 found to be approximately  $5.87 \times 10^7$  cells/ml.

#### 4.1: Preparation of feeder cells

Mouse peritoneal exudate cells were used as a feeder cell layer for hybrid cells. These were prepared the day 20 before the fusion was undertaken. Cold complete medium containing hypoxanthine, aminopterin and thymidine (HAT medium-see reagents) was used for preparation of the feeder cells. Five milliliters of the medium was injected into the peritoneal cavity of a freshly killed Balb/c mouse and after 25 gentle agitation was removed aseptically now containing exudate cells. One hundred microlitres of the cells were added to each well of a sterile 96 well plate (4-5 plates were needed per spleen to be fused). All plates were examined on the day of the fusion to ensure that they free from any 30 visible bacterial or fungal contamination.

#### 4.2: Preparation of spleen cells

Preparation of spleen cells from both MAM HBSAg immunised and the NP HBSAg immunised mice were carried out in 35 a similar method.

The spleen was carefully removed from the freshly killed mouse and placed in a petri dish containing 5mls of

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cold complete medium. The spleen was disrupted carefully with the blunt side of a scalpel blade thus teasing away the cells from the spleen connective tissue. The cell suspension was then transferred to a sterile universal container and the debris allowed to settle to the bottom. The supernatant was carefully removed and the cells pelleted at 150g for 10 minutes. The cells were washed three times in 10ml of incomplete medium and the final cell pellet was resuspended in 10ml of incomplete medium. The yield and viability of the spleen cells was determined using a Neubauer chamber. Approximately  $7 \times 10^7$  cells were harvested from the NP spleen whilst  $10 \times 10^7$  cells were obtained from the first MAM spleen and  $2.5 \times 10^7$  cells from the second MAM spleen.

#### 4.3: Cell fusion

The splenocytes were mixed with the myeloma cells at a 4:1 ratio and the mixture centrifuged at 1500g for 10 minutes. The supernatant was removed leaving a dry pellet of cells at the bottom the container. This was placed in a water bath at 37°C. Polyethylene glycol 1540 (PEG) which had been previously sterilised by autoclaving was also warmed in the 37°C waterbath. The cell pellet was loosened in the bottom of the container and 1ml of the PEG solution added dropwise whilst the container was gently agitated. The cell/PEG mix was left at 37°C for one minute and then an additional 2ml of the warm incomplete medium added. Further incomplete medium was added slowly so that the volume doubled every minute until 25ml of incomplete medium had been added. The fused cells were then centrifuged at 1500g for 10 minutes and the pellet resuspended in 10ml of HAT medium. The cell suspension was distributed evenly over five plates of feeder cells and then placed in a CO<sub>2</sub> incubator.

#### 4.4: Culture of hybrid cells

Clusters of growing fused cells appeared after about a week. At this stage the cells were re-fed by replacing 100 µl of old medium with fresh HAT medium. Approximately 14 days



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after the fusion, 100  $\mu$ l of the supernatant fluid from each well where successfully fused clusters of cells were found, was removed and assayed for the presence of anti-HBs. The supernatant was replaced with complete medium containing 5 hypoxanthine and thymidine (HT medium-see Reagents).

4.5: Reverse capture RIA for the detection of anti-HBs secreting hybridomas

Supernatant from viable hybridoma cultures were tested for anti-HBs on the basis of a reverse capture assay.

Round bottom Nunc microwells were coated with 100  $\mu$ l of a 1:1000 dilution in Tris buffer of the IgG fraction of rabbit anti-mouse IgG. After two days at room temperature the wells were washed with Tween Saline and quenched for one hour with 0.5% bovine serum albumin in Tris buffer (Tris BSA buffer-see reagents). The wells were then sealed and stored moist at 4°C. All washings were done with Tween Saline.

Before the assay was carried out, the Tris BSA buffer was removed. One hundred microlitres of a 1:10 dilution in phosphate buffered saline of the supernatant from each well where successfully fused clusters of cells were found was added to the assay wells and incubated at 37°C for one hour. After washing, 100  $\mu$ l of purified  $^{125}$ I-wild type HBsAg were added and left in a moist box at room temperature overnight. The assay wells were then washed and bound reactivity measured in a sixteen channel gammacounter. Supernatant containing anti-HBs gave an increased binding of the label. The hybridomas were subsequently retested by the same assay using  $^{125}$ I-MAM and  $^{125}$ I-NP labels instead of  $^{125}$ I-wild type HBsAg. Each assay carried out included positive and negative controls. Positive controls used were monoclonals D2H5 and H3F5 raised against wild type HBsAg. An anti-HIV gag monoclonal 3D3F2 and phosphate buffered saline were used as negative controls.

From the results of the assay, parent wells containing colonies secreting anti-HBs were chosen for further screening and cloning.

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When the cells were growing well, they were aspirated using a glass pasteur pipette and transferred aseptically to 24 well sterile plates which contained a layer of mouse peritoneal exudate cells (PEC) in HT medium.

5

#### 4.6: Cloning by limiting dilution

Hybridomas identified as anti-HBs positive were cloned by limiting dilution. This is done to ensure that the antibody secreted is homogenous and monospecific. For each  
10 hybrid culture the volume of cell suspension containing 100 cells was calculated and this volume added to 10ml of complete medium. One hundred microlitres of this cell suspension was added to each of 48 wells of a 96 well plate containing feeder cells in complete medium. One hundred more  
15 cells were added to the remaining 5ml of complete medium and 100 $\mu$ l of this more concentrated cell suspension were added to 24 wells of the plate. Finally, a further 100 cells were added to the remaining cell suspension and this was distributed over the last 24 wells. Variation in the  
20 concentration of cells on each plate made allowance for inaccurate cell counts and low viability of some cultures. This procedure was repeated for each selected parent hybridoma.

Approximately five days after cloning the plates were  
25 examined for wells where only a single colony was visible. One hundred microlitres of supernatant was removed from these wells and retested for anti-HBs using the reverse capture assay described previously.

Clones secreting anti-HBs were then expanded as before  
30 and re-fed as necessary by removal of 100 $\mu$ l of tissue culture supernatant and replaced with fresh complete medium.

#### 4.7: Culture of hybrid clones

Once the colonies were large enough to cover the well  
35 they were expanded into 12cm<sup>2</sup> tissue culture flasks containing feeder cells and then further into 25cm<sup>2</sup> and 75cm<sup>2</sup> flasks. The cells were split 1:2 approximately every 2-3 days

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with fresh complete medium.

Once good growth had been established, some of the cells were frozen and stored under liquid nitrogen in incomplete medium supplemented with 10% dimethyl sulphoxide and 50% foetal calf serum.

#### 4.8: Production of ascitic fluid

Female Balb/c mice, aged 12-20 weeks were primed with 0.5ml pristane. One to four weeks after priming, 1ml of cloned hybridoma cells, resuspended in incomplete medium, were injected intraperitoneally. The ascitic fluid was aspirated 1-3 weeks later and separated from cells by centrifugation at 3000g for 10 minutes. The ascitic fluid was then stored at -20°C.

15

## II. CHARACTERISATION OF THE MONOCLONAL ANTIBODIES RAISED AGAINST WILD TYPE AND MUTANT HBsAg

### 1: Isotyping of monoclonal antibodies

Isotyping of the monoclonal antibodies was carried out using a kit from Serotec (Serotech Limited, 22 Bankside Station Approach, Kidlington, Oxford OX5 1BR) specifically designed to identify the class and subclass of monoclonal antibodies in tissue culture supernatants. Used in accordance with the manufacturer's recommendations, it readily characterised the mouse monoclonal antibody.

### 2: Serum protein electrophoresis

Ascitic fluid from each clone was tested for the presence of a monoclonal protein band by serum protein electrophoresis (SPE, Paragon Electrophoresis System, Beckman Ltd). Following the protocol provided, the kit conveniently identified those ascitic fluids containing high levels of monoclonal immunoglobulin.

The Paragon Serum Protein Electrophoresis kit is intended for the electrophoretic separation of proteins in a buffered agarose gel. After electrophoresis, the proteins in the gel were immobilised in a fixative solution and the gel

dried to a film. The protein pattern is visualised by staining the film with a protein-specific stain and the pattern visually interpreted.

5 3: Immunoglobulin G preparation

Immunoglobulin G was prepared by ion exchange chromatography from the various ascitic fluids.

DE52 gel (Whatman Ltd), supplied pre-swollen was resuspended in a 0.2M phosphate buffer pH 8.0. The gel was  
10 then dispersed in distilled water to give a 10mM buffer strength.

A K9 column (Pharmacia) was packed with the DE52 gel to give a gel to sample volume ratio of 5:1. All columns were equilibrated with an equal volume of 10mM phosphate buffer  
15 (PB-see reagents). After overnight dialysis at 4°C in 10mM PB, the sample was layered on top of the columns and allowed to absorb over 30 minutes.

Immunoglobulin G from the ascitic fluid was recovered by stepwise elution with 10mM, 30mM and 60mM PB. Each buffer  
20 was allowed to run through the column and the optical density of the eluent was monitored and recorded at 280nm (Ultra Violet Spectrophotometre, LKB Ltd.). The three different eluents were collected separately and assayed for anti-HBs activity in the reverse capture RIA. The eluent containing  
25 the majority of anti-HBs reactivity could be identified.

The protein concentrations of the ascitic fluid IgG were calculated by determining their absorbance at 280nm using an E1% 1cm value of 1.4 in a spectrophotometer.

30 4: Immunoglobulin A preparation

As with IgG, IgA was prepared using ion exchange chromatography. Separation was, however, carried out on a Sephacryl gel. As with DE52, the gel was equilibrated with 10mM phosphate buffer. The sample was layered on top of the  
35 column and allowed to absorb over 30 minutes. As before, the ascitic fluid was separated by stepwise elution. The eluent was monitored and then tested for anti-HBs reactivity.

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### 5: Radiolabelling procedure

Labelling of the immunoglobulin fractions was carried out by the iodogen method (Salacinski et al., 1979).

Clean 7.5x10cm glass tubes were initially coated with 5  $\mu\text{g}$  of chloroform. To the iodogen tubes 15  $\mu\text{g}$  of protein in PBS was added. Finally  $\text{Na}^{125}\text{I}$  (0.5 mCi, Amersham International Plc) was added and the reaction in the tube allowed to proceed for 10 minutes on ice. The iodogen acts as a mild oxidising agent for the  $\text{Na}^{125}\text{I}$  to bind onto a tyrosine 10 residue on the protein.

A K9 column was then packed with Sephadex G-25 and equilibrated in Tris BSA buffer. Non-radioactive iodide (KI/NaI) in PBS was added to the G25 column as this reduced the tendency for free  $^{125}\text{I}$  to stick to the column. The 15 reaction mixture was then removed from the iodogen tube and transferred to the column. Tris BSA buffer was then used for eluting and the eluent monitored. The  $^{125}\text{I}$ -labelled protein fraction was collected from the first peak and stopped as the peak began to tail off. The labelled protein was then stored 20 at 4°C in Tris saline buffer containing 5% BSA.

Elution was continued until free  $^{125}\text{I}$  had come through (i.e. the second peak). The height of this peak compared to the height of the first peak gave an estimate of the percentage of  $^{125}\text{I}$  bound.

25

## III. RESULTS

### 1: Patients

The sera from both patients MAM and NP were tested in a series of assays for HBsAg. Results of the comparison of 30 monoclonal and polyclonal antibody based methods is shown in Table 1 below.

Table 1: HBsAg assay detection: a comparison

Patient	Assay 1	Assay 2	Assay 3	Assay 4	Assay 5
NP	+	-	+	+	+
MAM	+	-	+	nt	nt

Key to Table 1:

- Assay I Reverse passive haemagglutination assay.  
 Assay II Monoclonal based antibody assay.  
 Assay III Mixed monoclonal/polyclonal based assays.  
 Assay IV Mixed monoclonal/polyclonal based assays.  
 Assay V Monoclonal based antibody assay.

Results from the sequencing revealed several point mutations in the "a" determinant of both patients. Mutations were found in HBV DNA amplified from patient NP (subtype ayw) at surface antigen codons encoding amino acids 133, 134 and 144. Mutations were found at codons encoding amino acids 133, 134, 142, 144 and 145 in HBV DNA amplified from patient MAM (subtype adr). Those mutations were detected both before and after transplantation. The sequences did not differ between the period when HBsAg was detectable by either polyclonal antibody based assays alone as compared with the time when HBsAg was detectable by both monoclonal and polyclonal based assays. The mutations are shown in Table 2.

MAM HBsAg is Mutant HBsAg I and NPHBsAg is Mutant HBsAg II. The complete amino acid sequence of, and nucleotide sequence encoding, HBsAg is given in Valenzuela et al. (1979).

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Table 2: Mutations detected in the HBsAg "a" determinant of patients MAM and NP

5	Patient	Codon	Wild Type	Wild Type	Mutant	Mutant
			Codon	Amino Acid	Codon	Amino Acid
10	NP	133	ATG	Methionine	ATT	Isoleucine
		134	TAT	Phenylalanine	CAT	Histidine
		144	GAT	Aspartic Acid	GTC	Valine
15	MAM	133	ATG	Methionine	ATC	Isoleucine
		134	TAT	Phenylalanine	AAT	Asparagine
		142	CCT	Proline	AGT	Serine
		143	TCG	Serine	TTG	Leucine
		145	GGA	Glycine	AAA	Lysine
20	-----					

## 2: Identification and distribution of anti-HBs secreting monoclonal antibodies

Hybridomas secreting anti-HBs were initially identified by using the reverse capture assay (see section II, 4.5). Approximately a week after the fusion all wells containing clusters of growing cells were tested for anti-HBs activity. Every assay performed included both negative and positive controls.

Cells arising from the MAM fusion were tested in the RIA in the first instance with the  $^{125}\text{I}$  MAM HBsAg whilst those from the NP fusion were tested with the  $^{125}\text{I}$  NP HBsAg. Results of the assay would give an indication of those clones which were producing antibodies against their respective surface antigens. Hybridomas which gave counts above 750cpm were considered to be positive. It was shown that of the 320 MAM clones tested, 10% were positive whilst 8% of the 200 NP clones were found to be positive.

To check for any reactivity against the wild type (WT) HBsAg, clones from both the NP and MAM fusions, regardless of whether they were positive in the previous assay were tested in the reverse capture RIA using  $^{125}\text{I}$  WT HBsAg label. As before, a positive result was indicated by an increased binding of the label. Hybridomas which gave counts over

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650cpm were considered as positive. Of the 320 MAM clones tested, 6% gave positive results whilst 4% of the NP clones were found to bind to the  $^{125}\text{I}$  WT HBsAg.

With the results obtained thus far it was possible to  
5 identify those clones secreting anti-HBs which was specifically recognising and therefore binding either MAM, NP or WT hepatitis B surface antigen.

To determine if there was any cross reactivity within these hybridomas the NP clones were tested with  $^{125}\text{I}$  MAM  
10 HBsAg and the MAM clones with the  $^{125}\text{I}$  NP HBsAg. It was found that 4 of the MAM clones recognised NP label whilst 5 of the NP clones were recognised by the MAM label.

The clones had now been tested with all label combinations. It was therefore possible to identify those  
15 hybridomas which had consistently given strong positive results be it with the  $^{125}\text{I}$  MAM,  $^{125}\text{I}$  NP or  $^{125}\text{I}$  WT HBsAg. The monoclonal antibodies could now be divided into the following categories: 1) MAM specific; 2) NP specific; 3) MAM/NP specific; 4) MAM/WT specific; 5) NP/WT specific;  
20 6) NP/WT/MAM specific.

Eight NP and ten MAM parent hybridomas each of which fell under one of the categories listed above were chosen. These hybridomas were chosen as they displayed good cell growth as well as being repeatedly positive when tested for  
25 anti-HBs activity.

The eighteen chosen hybridomas were then cloned by limiting dilution. Approximately fourteen days after cloning, wells containing single colonies were tested in the reverse capture assay with all three of the labels. Six of the eight  
30 NP hybridomas cloned successfully and were found to be still positive for anti-HBs activity.

However, with the MAM clones, it was found that only six of the ten chosen hybridomas cloned. Attempts at recloning the hybridomas from their respective parent  
35 colonies proved unsuccessful despite increasing the concentration of the cells added to each plate. The six successfully cloned hybridomas containing single colonies



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were tested for anti-HBs activity. From the six, two hybridomas, M3C9 and M3A10, both three way crosses were found to be still positive for anti-HBs. Hybridomas derived from the MAM fusion were proving to be extremely unstable as they 5 were passaged.

At least three positive clones were chosen from each hybridoma for expansion. The clones were propagated in culture and retested two weeks later. All the NP clones were still positive for anti-HBs secretion. However of the MAM 10 clones, M3C9 had also proved to be extremely unstable giving negative results in the RIA.

The final positive secreting hybridomas from each fusion are shown in Table 3 below. The table also gives the monoclonal antibody specificity and their RIA values rounded 15 up to the nearest 500cpm.

**Table 3: Clones derived from NP and MAM fusions with their respective specificity and RIA values**

20	Clone	Binding of WT	$^{125}\text{I}$ HBsAg(cpm) MAM	NP	Clone specificity
25	*P2D3	4000	6000	3500	WT/NP/MAM cross
30	P2C6	neg	5000	3000	NP/MAM cross
35	P2H6	2000	neg	11000	NP/WT cross
	P2H9	neg	neg	1000	NP only
40	P4C11	neg	neg	10000	NP only

... continued

Table 3 continued

5	Clone	Binding of WT	<sup>125</sup> I MAM	HBsAg(cpm) NP	Clone specificity
10	P3E4	1000	neg	10000	NP/WT cross
15	*M3A10	3000	5000	2500	WT/NP/MAM cross
20	Control D2H5	7000	neg	2000	WT/NP cross
	Control H3F5	6000	1000	neg	WT/MAM cross

25 Where P represents the hybridomas derived from the NP fusion and M represents the hybridomas derived from the MAM fusion.

30 \*Denotes three-way cross.

35 A second MAM fusion was undertaken. The fusion was carried out as described before. However, the yield of spleen cells was only 25% of that recorded in the original MAM fusion. Approximately a week later, all wells containing clusters of growing cells were tested for anti-HBs activity using the reverse capture RIA. As before, each assay performed included negative and positive controls. All cells arising from the fusion were tested with each of the labels, <sup>125</sup>I MAM HBsAg, <sup>125</sup>I NP HBsAg and <sup>125</sup>I WT HBsAg.

40 Of the 68 cloned tested, only three produced positive results. One clone was found to be MAM specific whilst the other two, M4H2 and M4F5, were found to be WT/NP/MAM crosses (Table 4). However, as described in more detail below, M4H2 proved to be unstable, like some of the hybridomas from the first MAM fusion. It eventually stabilised as a MAM-specific clone.

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Table 4: Clones derived from second MAM fusion with their respective specificities and RIA values

Clone	Binding of WT	$^{125}\text{I}$ HBsAg (cpm) MAM	NP	Clone specificity
M4B12	neg	7000	neg	MAM only
**M4H2	9000	8000	1500	WT/NP/MAM cross
*M4F5	10000	9000	10000	WT/NP/MAM cross
Control D2H5	10000	neg	2000	WT/NP cross
Control H3F5	10000	7000	neg	WT/MAM cross

\* Denotes three-way cross

\*\* Initially found to be a three-way cross but was later found to be unstable and changed to a MAM-specific clone, see below.

All three of the hybridomas displayed good cell growth and were cloned by limiting dilution. Cloning on this occasion proved successful and all wells containing single colonies were tested in the reverse capture RIA with all three of the labels. The clones were still found to be positive for anti-HBs activity.

Five positive clones were chosen for expansion. After further growth in culture, the clones were screened and found to be still positive for anti-HBs production.

The final number of positive parent hybridomas and their antibody specificity and cross-reactivity obtained from the NP and the two MAM fusions are shown in Table 5.

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Table 5: Final number of clones with their  
respective specificities and RIA values.

5	Clone	Binding of WT	<sup>125</sup> I HBsAg (cpm) MAM	NP	Clone specificity
10	*P2D3	4000	6000	3500	WT/NP/MAM cross
15	P2C6	neg	5000	3000	NP/MAM cross
20	P2H6	2000	neg	11000	NP/WT cross
	P2H9	neg	neg	1000	NP only
25	P4C11	neg	neg	10000	NP only
30	P3E4	1000	neg	10000	NP/WT cross
	*M3A10	3000	5000	2500	WT/NP/MAM cross
35	M4B12	neg	7000	neg	MAM only
40	**M4H2	9000	8000	1500	WT/NP/MAM cross
45	*M4F5	10000	9000	10000	WT/NP/MAM cross
	Control D2H5	10000	neg	2000	WT/NP cross
50	Control H3F5	10000	7000	neg	WT/MAM cross

55 For key: see following page

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Key to table 5:

\* Denotes three-way cross

5

\*\* Initially found to be a three-way cross but was later found to be unstable and changed to a MAM-specific clone, see below.

10 Further cross-competition studies were carried out using antibodies P2D3, M4F5 and M3A10 with WT, NP and MAM HBsAg. Each of those antibodies cross-competed with each other, but did not cross-compete with any of the other monoclonal antibodies raised. At this stage M4H2 still  
15 appeared to be a three-way cross.

These results further indicate that the monoclonal antibodies of the invention bind to a distinct epitope.

One positive clone from each hybridoma parent was chosen and injected into a previously pristane primed Balb/c  
20 mouse. One to three weeks after the intraperitoneal inoculation, the ascitic fluid was harvested from the mice. All clones successfully produced ascitic tumours in the mice. The quantity of ascitic fluid obtained from each mouse varied from 1ml to 5ml.

25 More extensive cross-competition studies were carried out using various different antibodies, each with WT, MAM and NP HBsAg. Those tests were carried out according to the reverse capture RIA protocol given above with the following modifications: the microwells were coated with a goat  
30 polyclonal anti-HBs antibody. In each case 100  $\mu$ l of the antigen were added and incubated overnight, then 50  $\mu$ l each of  $^{125}$ I labelled and unlabelled monoclonal antibody were added and binding was determined. The results are presented in Tables 6a, 6b and 6c. In those Tables, the labelled  
35 antibodies are listed horizontally and the unlabelled antibodies are listed vertically.

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Table 6a: WT HBSAg (1:3K)

	P4C11	P3E4	P2D3	P2H6	P2H9	P2C6	M4F5	M4H2	M3A10	D2H5	H3F5
P4C11	-	-	-	+++	-	-	-	-	-	-	-
P3E4	-	+++	-	+++	-	-	-	-	-	-	-
P2D3	-	-	+++	+++	-	-	+++	-	+++	-	-
P2H6	-	-	-	+++	-	-	-	-	-	-	-
P2H9	-	-	-	+++	-	-	-	-	-	-	-
P2C6	-	-	-	+++	-	-	-	-	-	-	-
M4F5	-	-	+++	+	-	-	+++	-	+++	-	-
M4H2	-	-	-	+++	-	-	-	-	-	-	-
M3A10	-	-	+++	+++	-	-	+++	-	+++	-	-
D2H5	-	-	-	-	-	-	-	-	-	+++	-
H3F5	-	-	-	-	-	-	-	-	-	-	+++
percentage binding	0.4	1.0	4.8	0.4	0.1	0.1	14	0.1	0.3	37	23

+++ > 75% inhibition  
 ++ 50-75% inhibition  
 - < 50% inhibition

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Table 6b: MAM HBsAg (1.3k)

	P4C11	P3E4	P2D3	P2H6	P2H9	P2C6	M4F5	M4H2	M3A10	D2H5	H3F5
P4C11	-	-	-	-	-	-	-	-	-	-	-
P3E4	-	-	-	-	-	-	-	-	-	-	-
P2D3	-	-	+++	-	-	-	+++	-	++	-	++
P2H6	-	-	-	-	-	-	-	-	-	-	-
P2H9	-	-	-	-	-	-	-	-	-	-	-
P2C6	-	-	-	-	-	-	-	-	-	-	-
M4F5	-	-	+++	-	-	-	+++	-	++	-	-
M4H2	-	-	-	-	-	-	-	+++	-	-	++
M3A10	-	-	++	-	-	-	+++	-	++	-	-
D2H5	-	-	-	-	-	-	-	-	-	-	-
H3F5	-	-	-	-	-	-	-	+++	-	-	+++
Percentage binding	0.1	0.3	6.7	0.4	0.3	2.2	21	14	0.8	0.2	23

+++ > 75% inhibition  
 ++ 50-75% inhibition  
 - < 50% inhibition

Table 6c: NP HBSAg (1:150)

	P4C11	P3E4	P2D3	P2H6	P2H9	P2C6	M4F5	M4H2	M3A10	D2H5	H3F5
P4C11	+++	-	-	+++	-	-	-	++	-	-	-
P3E4	-	+++	-	-	+++	-	-	-	-	-	-
P2D3	-	-	+++	++	++	-	+++	-	-	-	-
P2H6	+++	-	-	+++	++	-	-	++	-	-	-
P2H9	++	-	-	++	++	-	-	++	-	-	-
P2C6	-	-	-	-	++	+++	-	++	-	-	-
M4F5	-	-	+++	-	++	-	+++	-	-	-	-
M4H2	-	-	-	-	++	-	-	-	-	-	-
M3A10	-	-	++	-	++	-	++	-	-	-	-
D2H5	-	-	-	-	-	-	-	-	-	+++	-
H3F5	-	-	-	-	-	-	-	-	-	-	-
Percentage binding	6	3	2	9	0.3	0.9	4	0.2	0.1	21	0.4

+++ > 75% inhibition  
 ++ 50-75% inhibition  
 - < 50% inhibition



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The results shown in Tables 6a, 6b and 6c confirm that the three-way cross antibodies M4F5 and P2D3 are binding to a common epitope.

M3A10 was found previously to be a three-way cross  
5 antibody (see Table 4). The fact that it is an IgA antibody (see Section III.4 below) may explain the poor binding results shown in Table 6a since this class of antibody (IgA) is difficult to purify.

As shown above, the hybridoma M4H2, which was produced  
10 in the second MAM fusion, was initially found to be a three-way cross (see Table 4). However, after further growth in culture, that hybridoma showed decreasing binding to  $^{125}\text{I}$  WT HBsAg and to  $^{125}\text{I}$  NP HBsAg. Retesting for anti-HBs activity against the WT and NP HBsAg over a period of time revealed  
15 negative results. However, M4H2 continued to recognise and bind strongly to the  $^{125}\text{I}$  MAM HBsAg. It appears therefore, that the clone specificity was initially unstable and with the passage of time has evolved to a MAM-specific antibody-secreting hybridoma.

20 Binding of  $^{125}\text{I}$ -labelled antibodies to various different HBsAgS was measured using a RIA in which HBsAg was immobilised to the solid phase (microwells) via a polyclonal goat anti-HBs coated on the microwells. The HBsAgS used were WT, MAM and NP HBsAg, which have been described above, SP  
25 HBsAg and SZ HBsAg. SP HBsAg, which is serotype adw, has a mutation in the codon for amino acid 143 leading to substitution of met for ser. In SZ HBsAg, serotype adr, a mutation leads to substitution of arg for gly at position 145. (SP HBsAg is Mutant HBsAg IV and SZ HBsAg is Mutant  
30 HBsAg III.) A pool of normal human sera that has been tested and found to be hepatitis B marker free (NHS) was used as control. This is an example of an immunoassay according to the present invention. The results are presented in Table 7 below.

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Table 7

HBSAg	% Binding of $^{125}\text{I}$ - anti mutant MAb			
	P2D3 (3 x crosses)	M4F5	M4H2 (MAM specific)	P4C11 (NP specific)
WT	12	15	0.6	0.1
MAM	13	20	31	0.2
NP	10	12	0.2	10
SP	23	26	0.5	0.4
SZ	25	30	0.4	0.5
NHS	0.3	0.2	0.4	0.3

nt = not tested

These results show that the three-way cross antibodies detect wild type HBsAg and also HBsAg from various different mutants, including mutant HBsAg different from that used for screening the antibodies. These results further confirm that the three-way cross antibodies are binding to an epitope common to the wild type protein and to mutant forms and demonstrate the utility of the antibodies in HBsAg assays. Using such antibodies it is possible to detect the presence of escape mutants.

### 3: Isotyping of monoclonal antibodies

Isotyping of the monoclonal antibodies was carried out using a kit from Serotec. Each isotyping reagent consisted of a purified rat monoclonal antibody specific for a single class-subclass of immunoglobulin, coupled to sheep red blood cells. The coupled rat antibodies recognised the heavy

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chain portion of the mouse immunoglobulin. The principle of the test system was based on red cell agglutination where a positive agglutinated result was produced when highly specific antibody recognised and bound to the particular isotype to which it is directed. The binding formed a lattice on the bottom of the microtitre plate well. A negative result was produced when the reagent cells were put into a supernatant containing a class of antibody which they did not recognise. The reagent cells fell to the bottom of the well forming a small "button".

The isotype of the monoclonal antibodies are shown in Table 8.

Table 8: Immunoglobulin classes of monoclonal antibodies

Monoclonal Antibody	Immunoglobulin Class
P4C11	IgG1
P3E4	IgG1
*P2D3	IgG1
P2H6	IgG1
P2H9	IgG2
P2C6	IgG1
*M3A10	IgA
**M4H2	IgG1

... continued

Table 8 continued

5		
	Monoclonal Antibody	Immunoglobulin Class
10	M4B12	IgG1
	*M4F5	IgG1
15		

\* Denotes three-way cross

20 \*\* Initially found to be a three-way cross but was later found to be unstable and changed to a MAM-specific clone, see below.

25 4: SPE analysis and purification of anti-HBs from ascitic fluid

Monoclonal protein was demonstrated in the ascitic fluid of all of the clones by serum protein electrophoresis. The electrophoretic mobility of the monoclonal protein band varied slightly for each different monoclonal ascitic fluid, but was identical for different batches of ascitic fluid obtained from the same clone. The disparity in migration distance between clones was a result of the differing ionic charges on the monoclonal proteins. The intensity of staining observed in the SPE monoclonal band gave an indication of the quantity of monoclonal protein present. This varied between ascitic fluid which had been harvested from different clones. A SPE band, however, was not seen with M4B12, the MAM specific monoclonal antibody. When the recovered ascitic fluid was tested in the RIA for anti-HBs activity, a negative result was given; all the other ascitic fluids gave the expected positive results. As before the MAM specific monoclonal antibody was proving to be unstable changing from a positive secreting hybridoma to a negative. (The M4B12 parent hybridoma has since been recloned and will be retested).

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Preparation of immunoglobulin from the ascitic fluids was carried out as described in sections III, 3 and III, 4 above. Immunoglobulin G was prepared by ion exchange chromatography on DE52 whilst IgA was separated on a Sephacryl column. Three separate protein peaks were obtained when increasing the ionic strength from 10mM to 30mM and finally 60mM PB for each monoclonal ascitic fluid. The size of the peaks varied from one monoclonal fluid to another. Eluent was collected at each peak. The anti-HBs IgG/IgA in the different buffer concentrations were then measured in the reverse capture assay with  $^{125}\text{I}$  MAM HBsAg,  $^{125}\text{I}$  NP HBsAg and  $^{125}\text{I}$  WT HBsAg. The assay allowed a direct comparison of results between different buffer fractions to be made. From the assay, it was possible to choose the fraction showing the highest anti-HBs activity. The protein concentrations of the chosen fractions were determined (Table 9). The quantity of IgG/IgA obtained from each monoclonal antibody varied between each batch of ascitic fluid harvested.

Table 9: Quantity of IgG/IgA harvested from 1ml of each monoclonal ascitic fluid

Monoclonal antibody	Quantity of IgG/IgA from 1ml of ascitic fluid (mg/ml-1)
P4C11	4.00
P3E4	4.40
*P2D3	1.14
P2H6	2.40
P2H9	0.90

... continued

Table 9 continued

5	Monoclonal antibody	Quantity of IgG/IgA from 1ml of ascitic fluid (mg/ml-1)
10	P2C6	3.00
15	*M4F5	1.00
20	**M4H2	1.70
	*M3A10	2.00

25 \* Denotes three-way cross

\*\* Initially found to be a three-way cross but was later found to be unstable and changed to a MAM-specific clone, see below.

30

The intensity of the monoclonal band staining in SPE showed strong positive correlation to the protein concentration results in Table 7. For example P4C11 showed a very darkly stained band in SPE and had a high protein concentration whilst P2H9 showed a faintly stained band and had a low protein concentration.

#### GENERAL DISCUSSION

40 The results of HBsAg detection assays have confirmed that as a result of mutations, HBsAg in serum may not be detected by some monoclonal antibody based systems. Such mutations may cause the failure of either the capture antibody or the conjugated detection antibody to bind. These results show the importance of testing for a second marker in suspected cases of HBV infections when the initial screening test for HBsAg is negative. Both NP and MAM would have shown

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an anti-HBc only profile if tested for HBsAg with a monoclonal-based assay. The presence of HBsAg in the samples was revealed on testing by polyclonal based systems.

A mutation at amino acid 145 (glycine to lysine) was detected in patient MAM. Although this mutation was different from that previously described in vaccine escape mutants (glycine to arginine), the amino acid substitution was such that the predicted effect on the antigen structure is similar.

10 Patient NP did not have a mutation at codon 145 but multiple mutations as with patient MAM. Which of the mutations is responsible for the loss of detectability has not been determined. Interestingly in patient MAM the nucleotide sequences did not vary with change in HBsAg  
15 specificity in the HBsAg detection assays. This may have been due to the emergence of wild type virus present at less than 15-20% of the total viral load, the limit of detection by direct sequence analysis.

The production of monoclonal antibodies against the  
20 mutant hepatitis B surface antigens was successful. The number of clones produced in the NP and initial MAM fusion and the relative distribution of the specificities of the resulting antibodies were favourable. In order to identify anti-HBs secreting hybridomas, a sensitive but quick  
25 screening assay was required. An IgG reverse capture radioimmunoassay was chosen. The assay, though simple to set up, is capable of a very low background and a large dynamic range. The relative binding by positive clones of the labelled wild type and mutant HBsAg were unambiguously high  
30 and consistent giving reliable results.

Screening the hybridomas was initially carried out testing those clones derived from the MAM fusion with  $^{125}\text{I}$  MAM HBsAg and the NP clones with  $^{125}\text{I}$  NP HBsAg. Results of this assay gave an indication of those clones specifically  
35 producing anti-HBs which were recognising and thus binding to the MAM HBsAg or the NP HBsAg. All clones produced as a result of the two fusions were then tested with  $^{125}\text{I}$  WT

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HBsAg. No single clone gave a positive reaction to the WT HBsAg only. Either MAM/WT crosses or NP/WT crosses were found.

It was therefore possible to identify those clones  
5 which were recognising the MAM HBsAg or the NP HBsAg. Those clones which were also binding to the NP and WT HBsAg were also identified. To test for any cross reactivity between the hybridomas the clones produced from the MAM fusion were tested with the  $^{125}\text{I}$  NP HBsAg and vice versa. As a result of  
10 the screening assays, it was possible to categorise the clones as is listed in the Results section.

Positive controls included in the assay were two known WT HBsAg monoclonal antibodies, H3F5 and D2H5 whilst negative controls were PBS and 3D3F2, an anti-HIV gag monoclonal. As  
15 expected when used in the  $^{125}\text{I}$  WT HBsAg RIA, both H3F5 and D2H5 gave strong positive results. However when tested against  $^{125}\text{I}$  MAM HBsAg only H3F5 gave a positive reading and was negative when tested with  $^{125}\text{I}$  NP HBsAg. D2H5 though produced a positive signal with  $^{125}\text{I}$  NP HBsAg but was very  
20 weak when tested against  $^{125}\text{I}$  MAM HBsAg. Mutations in the MAM HBsAg eliminated the epitope for D2H5 whilst those in the NP HBsAg eliminated the H3F5 epitope.

The results obtained on testing with the SP and SZ mutant HBsAg show that the three-way cross antibodies are  
25 capable of detecting other mutant forms of HBsAg in addition to those used in the screening process. This confirms that the antibodies are binding to an epitope that is conserved between the wild type protein and mutant forms, and thereby confirms the value of including such antibodies in an HBsAg  
30 assay.

Although some of the hybridomas from the MAM fusions proved to be unstable, successful three-way crosses were obtained readily from both the MAM and NP fusions. Mutant forms of HBsAg other than the MAM and NP forms may be used  
35 for raising and/or for screening monoclonal antibodies according to the protocols described above.



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Note: Figure 1b is a typescript version of Figure 1a. In the case of any discrepancy between Figure 1a and Figure 1b, Figure 1a shall be taken as the authentic version.

## 5 Reagents

### Complete Medium

RPMI supplemented with

5mM	Hepes buffer
10 2Mm	L-glutamine
0.05M	2-Mercaptoethanol
20% v/v	fetal calf serum
25µg/ml <sup>-1</sup>	Fungizone
102 units ml <sup>-1</sup>	Penicillin
15 100µg ml <sup>-1</sup>	Streptomycin

### HAT medium

Complete medium plus

5x10 <sup>-5</sup> M	hypoxanthine
20 2x10 <sup>-3</sup> M	aminopterin
8x10 <sup>-4</sup> M	thymidine

### Incomplete Medium

RPMA supplemented with 5mM Hepes buffer

2nM	L-glutamine
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### HT medium

Complete medium plus

5x10 <sup>-3</sup>	hypoxanthine
8x10 <sup>-4</sup>	thymidine

### 0.02M Tris buffer pH7.6 (Tris buffer)

Sodium azide	1.00g
25 Tris (hydroxymethyl) methylamine	2.42g
Distilled water	900ml

Adjust pH to 7.6 with concentrated HCl and make that buffer up to 1 litre.

### 30 0.02M Tris BSA buffer

Add 0.5% BSA to 0.02M Tris buffer pH7.6.

### 0.2M Phosphate buffer (PB)

Solution A = 0.2M KH<sub>2</sub>PO<sub>4</sub>

35 Solution B = 0.2M Na<sub>2</sub>HPO<sub>4</sub>

Take 200ml solution B. Adjust pH to 8.0 with solution A.

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10, 20, 30 and 60 mM PB

For 10mM PB dilute 0.2M PB 1:20 in distilled water

For 30mM PB dilute 0.2M PB 1:6.6 in distilled water

For 60mM PB dilute 0.2M PB 1:3.3 in distilled water

5

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## CLAIMS:

1. A monoclonal antibody that is capable of binding specifically to wild-type HBsAg and to at least two mutant forms of HBsAg.
2. A monoclonal antibody as claimed in claim 1, wherein a mutant form of HBsAg has at least one amino acid substitution relative to wild type HBsAg.
3. A monoclonal antibody as claimed in claim 1 or claim 2, wherein a mutant form of HBsAg has the sequence of HBsAg present in an HBV escape mutant.
4. A monoclonal antibody as claimed in any one of claims 1 to 3, wherein a mutant form of HBsAg has at least one amino acid substitution in the "a" determinant or in the region of the "a" determinant.
5. A monoclonal antibody as claimed in any one of claims 2 to 4, wherein an amino acid substitution results from a point mutation.
6. A monoclonal antibody as claimed in any one of claims 1 to 5, wherein a mutant HBsAg has an amino acid substitution within the sequence encoding amino acids 133 to 145 of HBsAg.
7. A monoclonal antibody as claimed in any one of claims 1 to 6, wherein a mutant HBsAg has an amino acid substitution relative to wild type HBsAg at any one or more of positions 133, 134, 141, 142, 143, 144 and 145.
8. A monoclonal antibody as claimed in claim 7, wherein the substitution is at any one or more of positions 143, 144 and 145.
9. A monoclonal antibody as claimed in claim 7, wherein the substitution is in addition to one or more other substitutions in the "a" determinant or in another region.
10. A monoclonal antibody as claimed in claim 1, which is capable of binding specifically to wild type HBsAg and to at least one of the following mutant forms of HBsAg:  
Mutant HBsAg I ("NP" HBsAg): met to ile at amino acid 133; phe to his at amino acid 134; and asp to val at amino acid 144;  
Mutant HBsAg II ("MAM" HBsAg): met to ile at amino acid 133;

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phe to asn at amino acid 134; pro to ser at amino acid 142; ser to leu at amino acid 143; and gly to lys at amino acid 145;

Mutant HBsAg III ("SZ" HBsAg): gly to arg at amino acid 145;

Mutant HBsAg IV ("SP" HBsAg): ser to met at amino acid 143.

11. A monoclonal antibody that is capable of binding specifically to wild-type HBsAg and to at least one mutant HBsAg carrying an "a" determinant coded for by sequences having point mutations at any one or more of the codons encoding amino acids 143, 144 and 145.
12. A monoclonal antibody as claimed in any one of claims 1 to 11 that is an IgG, IgM or IgA immunoglobulin.
13. Monoclonal antibody P2D3 as produced by the hybridoma designated P2D3 and deposited at the ECACC under accession number ECACC 97042331.
14. Monoclonal antibody M3A10 as produced by the hybridoma designated M3A10 and deposited at the ECACC under accession number ECACC 97042330.
15. Monoclonal antibody M4F5 as produced by the hybridoma designated M4F5 and deposited at the ECACC under accession number ECACC 97042519.
16. A monoclonal antibody as claimed in any one of claims 1 to 15, in a humanised form.
17. A fragment or a derivative of a monoclonal antibody as claimed in any one of claims 1 to 16.
18. A method of producing a hybridoma capable of producing a monoclonal antibody as claimed in claim 1, which comprises immunising an animal with wild type HBsAg or a mutant form of HBsAg antigen, optionally in the form of an appropriate antigenic fragment or derivative thereof, immortalising antibody producing cells to form hybridomas, screening the hybridoma cultures against wild type HBsAg and two or more mutant forms of HBsAg, optionally in the form of an appropriate antigenic fragment or derivative thereof, and selecting those hybridomas that produce antibodies that bind to wild type HBsAg and two or more mutant forms of HBsAg.
19. A method as claimed in claim 18, wherein a mutant form

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of HBsAg is as defined in any one of claims 2 to 11.

20. A hybridoma capable of producing a monoclonal antibody as claimed in any one of claims 1 to 11.

21. Hybridoma designated P2D3 and deposited at the ECACC under accession number ECACC 97042331.

22. Hybridoma designated M3A10 and deposited at the ECACC under accession number ECACC 97042330.

23. Hybridoma designated M4F5 and deposited at the ECACC under accession number ECACC 97042519.

24. A method of producing a monoclonal antibody as claimed in claim 1, which comprises culturing a hybridoma as claimed in any one of claims 20 to 23 or obtainable by a method as claimed in claim 18 or claim 19 in vitro or in vivo, and obtaining the monoclonal antibody from the culture medium.

25. A monoclonal antibody obtainable by the method claimed in claim 24.

26. An immunoassay for the detection of HBsAg, which comprises contacting a sample under investigation with a monoclonal antibody as claimed in any one of claims 1 to 16 or claim 25, a fragment or derivative thereof as claimed in claim 17, or a combination of two or more thereof, and detecting any resulting antigen-antibody complex.

27. An immunoassay as claimed in claim 26, wherein the monoclonal antibody, fragment, derivative or combination thereof is used in combination with one or more other antibodies selected from polyclonal anti-HBs antibodies and other monoclonal anti-HBs antibodies.

28. An immunoassay as claimed in claim 26 or claim 27, in a homogeneous or heterogeneous format.

29. An immunoassay as claimed in claim 28, in a capture or a competitive format.

30. An immunoassay as claimed in any one of claims 26 to 29, wherein an immunoassay for the detection of antibodies to hepatitis B core protein (HBc) is carried out simultaneously with the assay for HBsAg.

31. An immunoassay kit that comprises a monoclonal antibody as claimed in any one of claims 1 to 16 or claim 25, a

- 54 -

fragment or derivative thereof as claimed in claim 17, or a combination of two or more thereof, and other reagents required for carrying out an immunoassay for HBsAg and optionally also reagents for detecting anti-HBc antibodies.

32. A kit as claimed in claim 31, which also comprises one or more other anti-HBs antibodies selected from polyclonal anti-HBs antibodies and other monoclonal anti-HBs antibodies.

33. A kit as claimed in claim 31 or claim 32, wherein the other reagents are selected from washing solutions, diluents, standard solutions, control reagents and labelled anti-HBs antibodies.

34. A solid phase suitable for use in an immunoassay on which is immobilised a monoclonal antibody as claimed in any one of claims 1 to 16 or claim 25, a fragment or derivative thereof as claimed in claim 17, or a combination of two or more thereof.

35. A solid phase as claimed in claim 34, wherein one or more further anti-HBs antibodies selected from polyclonal anti-HBs antibodies and other monoclonal anti-HBs antibodies are also immobilised on the solid phase.

36. A solid phase as claimed in claim 34 or claim 35, on which is also immobilised an agent capable of capturing anti-HBc antibodies.

37. An antiserum suitable for use therapeutically or prophylactically for passive immunisation which comprises a monoclonal antibody as claimed in any one of claims 1 to 16 or claim 25, a fragment or derivative as claimed in claim 17, or a combination of two or more thereof.

38. A composition suitable for use therapeutically or prophylactically for passive immunisation against HBV which comprises a monoclonal antibody as claimed in any one of claims 1 to 16 or claim 25, a fragment or derivative as claimed in claim 17, or a combination of two or more thereof, in admixture with a pharmaceutically suitable carrier.

39. An antiserum as claimed in claim 37 or a composition as claimed in claim 38, which also comprises one or more other antibodies selected from polyclonal anti-HBs antibodies and



- 55 -

other monoclonal anti-HBs antibodies.

40. A method of therapeutic or prophylactic passive immunisation against HBV infection, which comprises administering to a subject a therapeutically or prophylactically effective amount of a monoclonal antibody as claimed in any one of claims 1 to 16 or claim 25, a fragment or derivative as claimed in claim 17, or a combination of two or more thereof.

41. A method as claimed in claim 40, wherein one or more other antibodies selected from monoclonal and polyclonal anti-HBs antibodies are also administered.

42. An anti-idiotypic antibody to a monoclonal antibody as claimed in any one of claims 1 to 16 or claim 25, or a fragment or derivative thereof as claimed in claim 17.

43. Isolated Mutant HBsAg I, Mutant HBsAg II, Mutant HBsAg III or Mutant HBsAg IV as defined in claim 10.

44. A mutant HBsAg as claimed in claim 43, obtainable from a natural source.

45. A mutant HBsAg as claimed in 43, obtainable recombinantly.

46. A mutant HBsAg as claimed in claim 45, being glycosylated.

47. A fragment or derivative of a mutant HBsAg as claimed in any one of claims 43 to 46, said fragment or derivative being capable of binding specifically to a monoclonal antibody as claimed in any one of claims 1 to 16 or claim 25 or to a fragment or derivative as claimed in claim 17.

48. A fragment or derivative as claimed in claim 47, said fragment or derivative being immunogenic.

49. A vaccine composition that comprises a mutant HBsAg as claimed in any one of claims 43 to 46 or a fragment or derivative as claimed in claim 48, or a combination of two or more thereof, and a pharmaceutically suitable carrier.

50. A vaccine composition as claimed in claim 49, which further comprises one or more other immunogenic peptides or polypeptides.

51. A vaccine composition as claimed in claim 50, wherein

- 56 -

the immunogenic peptides or polypeptides are immunogenic HBsAg peptides or polypeptides.

52. A method of vaccinating a subject against HBV, which comprises administering to the subject a therapeutically effective amount of a mutant HBsAg as claimed in any one of claims 43 to 46, a fragment or derivative as claimed in claim 48, or a combination of two or more thereof, optionally in combination with one or more other immunogenic HBsAg peptides or polypeptides.

53. An isolated nucleic acid that encodes Mutant HBsAg I, Mutant HBsAg II, Mutant HBsAg III or Mutant HBsAg IV as defined in claim 10.

54. A nucleic acid that encodes a fragment or derivative of a mutant HBsAg as defined in claim 10, said fragment or derivative being capable of binding specifically to a monoclonal antibody as claimed in any one of claims 1 to 16 or claim 25 or to a fragment or derivative thereof as claimed in claim 17.

55. A nucleic acid as claimed in claim 54, wherein the fragment or derivative of the mutant HBsAg is immunogenic.

56. A composition that comprises a nucleic acid as claimed in any one of claims 53 to 55.

57. A vaccine composition that comprises a nucleic acid as claimed in claim 53 or claim 55 in a form suitable for expression of the encoded amino acid sequence when administered to a mammal.

58. A method of vaccination against HBV, which comprises administering to a subject a therapeutically or prophylactically effective amount of a nucleic acid as claimed in claim 53 or claim 57 or a vaccine composition as claimed in claim 57.

FIGURE 1a

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2/2

	codon 122			thr asn		
1)						
2)	leuleuproglyserthrthrserthrthrglyprocysargthrcysthrthrproalaglnglylesermetpheproser					
3)	ctaattccaggatcatcaaccaccagcacgggaccctgcagaaacctgcacgactcctgctcaagggaatctctatgtatccctcc					
4)	c	a	a	t	g	a
	ca	t				
	t		gt	a	a	
5)						
6)	t	a	ta	t	c	tc

	codon 143 145			codon 160		
2)	cyscyscysthrlysproseraspglyasncyscyscysthrleproilleproserthrtrpalapheglylyspheleutrpleu					
3)	tgttgctgtacaaaaccttcggatggaaactgcacacctgtattcccatcccatcctcctggttcggaataatcctatgggag					
4)	c	a	c	t	t	g
5)	c	agt	t	aa	t	cg
6)	c		tc		c	t

FIGURE 1b

420 Rec'd PCT/PTO 0 1 OCT 1999

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## PCT

## INTERNATIONAL SEARCH REPORT

(PCT Article 18 and Rules 43 and 44)

Applicant's or agent's file reference <b>JSvn/556 WO</b>	<b>FOR FURTHER ACTION</b> see Notification of Transmittal of International Search Report (Form PCT/ISA/220) as well as, where applicable, item 5 below.	
International application No. <b>PCT/GB 97/ 01161</b>	International filing date (day/month/year) <b>25/04/1997</b>	(Earliest) Priority Date (day/month/year) <b>25/04/1996</b>
Applicant <b>MUREX DIAGNOSTICS CORPORATION et al.</b>		

This International Search Report has been prepared by this International Searching Authority and is transmitted to the applicant according to Article 18. A copy is being transmitted to the International Bureau.

This International Search Report consists of a total of 5 sheets.

☒ It is also accompanied by a copy of each prior art document cited in this report.

1. ☒ Certain claims were found unsearchable (see Box I).
2. ☐ Unity of invention is lacking (see Box II).
3. ☒ The international application contains disclosure of a nucleotide and/or amino acid sequence listing and the international search was carried out on the basis of the sequence listing
  - ☐ filed with the international application.
  - ☒ furnished by the applicant separately from the international application,
    - ☐ but not accompanied by a statement to the effect that it did not include matter going beyond the disclosure in the international application as filed.
  - ☐ Transcribed by this Authority
4. With regard to the title,
  - ☒ the text is approved as submitted by the applicant.
  - ☐ the text has been established by this Authority to read as follows:
5. With regard to the abstract,
  - ☒ the text is approved as submitted by the applicant.
  - ☐ the text has been established, according to Rule 38.2(b), by this Authority as it appears in Box III. The applicant may, within one month from the date of mailing of this International Search Report, submit comments to this Authority.
6. The figure of the drawings to be published with the abstract is:  
Figure No. \_\_\_\_\_
  - ☐ as suggested by the applicant.
  - ☐ because the applicant failed to suggest a figure.
  - ☐ because this figure better characterizes the invention.

☒ None of the figures.

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## INTERNATIONAL SEARCH REPORT

Inter Application No  
PCT/GB 97/01161

## A. CLASSIFICATION OF SUBJECT MATTER

IPC 6 C12N15/36 C07K16/08 C12N15/06 C12N5/20 C12P21/08  
G01N33/569 G01N33/577 B01J20/00 A61K39/42 C07K16/42  
C07K14/02 A61K39/29 A61K48/00

According to International Patent Classification (IPC) or to both national classification and IPC

## B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC 6 C07K

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

## C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	WO 95 21189 A (IMPERIAL COLLEGE OF SCIENCE, TECHNOLOGY & MEDICINE) 10 August 1995  see table 2 see figure 5	1,3,12, 26,28, 29,31, 33-35
X	--- J. WATERS ET AL.: "Loss of the common "A" determinant of hepatitis B surface antigen by a vaccine-induced escape mutant." THE JOURNAL OF CLINICAL INVESTIGATION, vol. 90, no. 6, December 1992, pages 2543-2547, XP002041202 see abstract see page 2546, left-hand column, line 23 - line 24 --- -/--	11,12, 17,20, 26,28, 29,31-33

☒ Further documents are listed in the continuation of box C.

☒ Patent family members are listed in annex.

## \* Special categories of cited documents:

- \*A\* document defining the general state of the art which is not considered to be of particular relevance
- \*E\* earlier document but published on or after the international filing date
- \*L\* document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)
- \*O\* document referring to an oral disclosure, use, exhibition or other means
- \*P\* document published prior to the international filing date but later than the priority date claimed

- \*T\* later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
- \*X\* document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
- \*Y\* document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art.
- \*&\* document member of the same patent family

Date of the actual completion of the international search

19 September 1997

Date of mailing of the international search report

01.10.97

Name and mailing address of the ISA

European Patent Office, P.B. 5818 Patentlaan 2  
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Fax: (+ 31-70) 340-3016

Authorized officer

Nooij, F

# INTERNATIONAL SEARCH REPORT

International Application No  
PCT/GB 97/01161

## C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	<p>WO 94 21812 A (IMPERIAL COLLEGE OF SCIENCE, TECHNOLOGY &amp; MEDICINE) 29 September 1994</p> <p>see the whole document</p> <p style="text-align: center;">---</p>	<p>11,12, 16,17, 26,28, 29,31-33</p>
A	<p>W. CARMAN ET AL.: "Fulminant reactivation of hepatitis B due to envelope protein mutant that escaped detection by monoclonal HBsAg ELISA." THE LANCET, vol. 345, no. 8962, 3 June 1995, LONDON, GB, pages 1406-1407, XP002041203 see the whole document</p> <p style="text-align: center;">---</p>	<p>1-36</p>
P,X	<p>X. QIU ET AL.: "Identification and characterization of a C(K/R)TC motif as a common epitope present in all subtypes of hepatitis B surface antigen." THE JOURNAL OF IMMUNOLOGY, vol. 156, no. 9, 1 May 1996, BALTIMORE, MD, USA, pages 3350-3356, XP002041204 see page 3352, right-hand column, line 7 - page 3353, left-hand column, line 7</p> <p style="text-align: center;">-----</p>	<p>1,2,4,5, 12, 18-20, 24,26, 28,29, 31,33</p>

# INTERNATIONAL SEARCH REPORT

International application No.  
PCT/GB 97/01161

## Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)

This International Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☒ Claims Nos.:  
because they relate to subject matter not required to be searched by this Authority, namely:  
see FURTHER INFORMATION sheet PCT/ISA/210
2. ☐ Claims Nos.:  
because they relate to parts of the International Application that do not comply with the prescribed requirements to such an extent that no meaningful International Search can be carried out, specifically:
3. ☐ Claims Nos.:  
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

## Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

1. ☐ As all required additional search fees were timely paid by the applicant, this International Search Report covers all searchable claims.
2. ☐ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. ☐ As only some of the required additional search fees were timely paid by the applicant, this International Search Report covers only those claims for which fees were paid, specifically claims Nos.:
4. ☐ No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Remark on Protest

- ☐ The additional search fees were accompanied by the applicant's protest.
- ☐ No protest accompanied the payment of additional search fees.

FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

Remark : Although claims 40,41,52 and 58 are directed to a method of treatment of the human/animal body , the search has been carried out and based on the alleged effects of the compound/composition.

# INTERNATIONAL SEARCH REPORT

Information on patent family members

International Application No

PCT/GB 97/01161

Patent document cited in search report	Publication date	Patent family member(s)	Publication date
WO 9521189 A	10-08-95	AU 1542695 A	21-08-95
		CN 1143373 A	19-02-97
		EP 0741745 A	13-11-96
		PL 315784 A	09-12-96
-----			
WO 9421812 A	29-09-94	AU 6287094 A	11-10-94
		CA 2158866 A	29-09-94
		CN 1121357 A	24-04-96
		EP 0689605 A	03-01-96
		JP 8510635 T	12-11-96
		NO 953749 A	22-11-95
		PL 310699 A	27-12-95
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## NOTES TO FORM PCT/ISA/220 (continued)

The letter must indicate the differences between the claims as filed and the claims as amended. It must, in particular, indicate, in connection with each claim appearing in the international application (it being understood that identical indications concerning several claims may be grouped), whether

- (i) the claim is unchanged;
- (ii) the claim is cancelled;
- (iii) the claim is new;
- (iv) the claim replaces one or more claims as filed;
- (v) the claim is the result of the division of a claim as filed.

The following examples illustrate the manner in which amendments must be explained in the accompanying letter:

1. [Where originally there were 48 claims and after amendment of some claims there are 51]:  
"Claims 1 to 29, 31, 32, 34, 35, 37 to 48 replaced by amended claims bearing the same numbers; claims 30, 33 and 36 unchanged; new claims 49 to 51 added."
2. [Where originally there were 15 claims and after amendment of all claims there are 11]:  
"Claims 1 to 15 replaced by amended claims 1 to 11."
3. [Where originally there were 14 claims and the amendments consist in cancelling some claims and in adding new claims]:  
"Claims 1 to 6 and 14 unchanged; claims 7 to 13 cancelled; new claims 15, 16 and 17 added." or  
"Claims 7 to 13 cancelled; new claims 15, 16 and 17 added; all other claims unchanged."
4. [Where various kinds of amendments are made]:  
"Claims 1-10 unchanged; claims 11 to 13, 18 and 19 cancelled; claims 14, 15 and 16 replaced by amended claim 14; claim 17 subdivided into amended claims 15, 16 and 17; new claims 20 and 21 added."

### "Statement under article 19(1)" (Rule 46.4)

The amendments may be accompanied by a statement explaining the amendments and indicating any impact that such amendments might have on the description and the drawings (which cannot be amended under Article 19(1)).

The statement will be published with the international application and the amended claims.

**It must be in the language in which the international application is to be published.**

It must be brief, not exceeding 500 words if in English or if translated into English.

It should not be confused with and does not replace the letter indicating the differences between the claims as filed and as amended. It must be filed on a separate sheet and must be identified as such by a heading, preferably by using the words "Statement under Article 19(1)."

It may not contain any disparaging comments on the international search report or the relevance of citations contained in that report. Reference to citations, relevant to a given claim, contained in the international search report may be made only in connection with an amendment of that claim.

### Consequence if a demand for international preliminary examination has already been filed

If, at the time of filing any amendments under Article 19, a demand for international preliminary examination has already been submitted, the applicant must preferably, at the same time of filing the amendments with the International Bureau, also file a copy of such amendments with the International Preliminary Examining Authority (see Rule 62.2(a), first sentence).

### Consequence with regard to translation of the international application for entry into the national phase

The applicant's attention is drawn to the fact that, where upon entry into the national phase, a translation of the claims as amended under Article 19 may have to be furnished to the designated/elected Offices, instead of, or in addition to, the translation of the claims as filed.

For further details on the requirements of each designated/elected Office, see Volume II of the PCT Applicant's Guide.

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## PACT COOPERATION TREATY

PCT

## NOTIFICATION OF ELECTION

(PCT Rule 61.2)

From the INTERNATIONAL BUREAU

To:

United States Patent and Trademark  
Office  
(Box PCT)  
Crystal Plaza 2  
Washington, DC 20231  
ETATS-UNIS D'AMERIQUE

in its capacity as elected Office

<b>Date of mailing</b> (day/month/year) 11 December 1997 (11.12.97)	
<b>International application No.</b> PCT/GB97/01161	<b>Applicant's or agent's file reference</b> JSvn/556 WO
<b>International filing date</b> (day/month/year) 25 April 1997 (25.04.97)	<b>Priority date</b> (day/month/year) 25 April 1996 (25.04.96)
<b>Applicant</b> TEDDER, Richard, Seton et al	

1. The designated Office is hereby notified of its election made:

☒ in the demand filed with the International Preliminary Examining Authority on:  
24 November 1997 (24.11.97)

☐ in a notice effecting later election filed with the International Bureau on:

2. The election ☒ was  
☐ was not

made before the expiration of 19 months from the priority date or, where Rule 32 applies, within the time limit under Rule 32.2(b).

The International Bureau of WIPO  
34, chemin des Colombettes  
1211 Geneva 20, Switzerland

Facsimile No.: (41-22) 740.14.35

Authorized officer

S. De Michiel

Telephone No.: (41-22) 338.83.38

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# PATENT COOPERATION TREATY



## PCT

### INTERNATIONAL PRELIMINARY EXAMINATION REPORT

(PCT Article 36 and Rule 70)

Applicant's or agent's file reference JSvn/556 WO		<b>FOR FURTHER ACTION</b>	See Notification of Transmittal of International Preliminary Examination Report (PCT/IPEA/416)
International application No. PCT/GB97/01161	International filing date (day/month/year) 25/04/1997	Priority date (day/month/year) 25/04/1996	
International Patent Classification (IPC) or national classification and IPC C12N15/36			
Applicant MUREX DIAGNOSTICS CORPORATION et al.			

<p>1. This international preliminary examination report has been prepared by this International Preliminary Examining Authority and is transmitted to the applicant according to Article 36.</p> <p>2. This REPORT consists of a total of 5 sheets, including this cover sheet.</p> <p><input type="checkbox"/> This report is also accompanied by ANNEXES, i.e., sheets of the description, claims and/or drawings which have been amended and are the basis for this report and/or sheets containing rectifications made before this Authority (see Rule 70.16 and Section 607 of the Administrative Instructions under the PCT).</p> <p>These annexes consist of a total of      sheets.</p>
<p>3. This report contains indications relating to the following items:</p> <ul style="list-style-type: none"> <li>I    <input checked="" type="checkbox"/> Basis of the report</li> <li>II   <input type="checkbox"/> Priority</li> <li>III <input type="checkbox"/> Non-establishment of opinion with regard to novelty, inventive step and industrial applicability</li> <li>IV <input type="checkbox"/> Lack of unity of invention</li> <li>V    <input checked="" type="checkbox"/> Reasoned statement under Article 35(2) with regard to novelty, inventive step or industrial applicability; citations and explanations supporting such statement</li> <li>VI <input type="checkbox"/> Certain documents cited</li> <li>VII <input type="checkbox"/> Certain defects in the international application</li> <li>VIII <input type="checkbox"/> Certain observations on the international application</li> </ul>

Date of submission of the demand 24/11/1997	Date of completion of this report 15.07.98
Name and mailing address of the IPEA/   European Patent Office D-80298 Munich Tel. (+49-89) 2399-0, Tx: 523656 epmu d Fax: (+49-89) 2399-4465	Authorized officer  Heimann-Pohl, B  Telephone No. (+49-89) 2399-8713 

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**INTERNATIONAL PRELIMINARY  
EXAMINATION REPORT**

International application No. PCT/GB97/01161

**I. Basis of the report**

1. This report has been drawn on the basis of (*substitute sheets which have been furnished to the receiving Office in response to an invitation under Article 14 are referred to in this report as "originally filed" and are not annexed to the report since they do not contain amendments.*):

**Description, pages:**

1-50 as originally filed

**Claims, No.:**

1-58 as originally filed

**Drawings, sheets:**

1/2-2/2 as originally filed

2. The amendments have resulted in the cancellation of:

- ☐ the description, pages:  
☐ the claims, Nos.:  
☐ the drawings, sheets:

3. ☐ This report has been established as if (some of) the amendments had not been made, since they have been considered to go beyond the disclosure as filed (Rule 70.2(c)):

4. Additional observations, if necessary:

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**INTERNATIONAL PRELIMINARY  
EXAMINATION REPORT**

International application No. PCT/GB97/01161

**V. Reasoned statement under Article 35(2) with regard to novelty, inventive step or industrial applicability; citations and explanations supporting such statement**

**1. Statement**

Novelty (N)	Yes:	Claims	9, 13-15, 18, 19, 21-24, 27, 30, 36, 37, 39-42, 47, 48, 50-58
	No:	Claims	1-8, 10-12, 16, 17, 20, 26, 28, 29, 31-35, 38, 43-46, 49
Inventive step (IS)	Yes:	Claims	13-15, 21-23
	No:	Claims	9, 18, 19, 24, 25, 27, 30, 36, 37, 39-42, 47, 48, 51, 58
Industrial applicability (IA)	Yes:	Claims	1-39, 41-51, 53-57
	No:	Claims	

**2. Citations and explanations**

**see separate sheet**

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- 1). The present application relates to monoclonal antibodies capable of binding to wild-type (wt) HBsAg and to at least two mutant forms of HBsAg (claims 1-10) and further to a monoclonal antibody that is capable of binding to wt HBsAg and to at least one mutant HBsAg carrying an "a" determinant coded for by sequences having point mutations at any one or more codons encoding amino acids 143, 144 and 145 (claim 11). The application is further directed to hybridomas and mutant HBsAg and their uses in diagnosis and prevention of HBV infections.
- 2). D1 (WO-A-95 21189) discloses, i.a., the monoclonal antibody RFHBs-2 which recognizes wt HBsAg and a mutant form of HBsAg in which at least two amino acids are inserted downstream of position 122 of the wt HBsAg. D1 further discloses assays and uses of said antibody and also the escape mutant of HBsAg with a gly to arg substitution at position 145.  
  
D2 (The Journal of Clinical Investigation, vol. 90, no. 6, 1992, pages 2543-2547) discloses further features of the monoclonal antibody RFHBs-2, namely, that it binds also to an escape mutant of HBsAg having a gly to arg substitution at position 145.  
  
D3 (WO-A-94 21812) discloses monoclonal antibodies raised against the escape mutant of HBsAg with a gly to arg substitution at position 145. Two of the antibodies, SMH HBs 145/G/R/I and SMH HBs 145/G/R/II, bind both to the wt and the variant.
- 3). Binding to wt and at least two mutant forms of HBsAg is a property of monoclonal antibody RFHBs-2 disclosed in D1 and D2 and one mutant form has a substitution in the "a" region. As a consequence the monoclonal antibodies of claims 1-8, 10-12 lack novelty (Art. 33 (2) PCT).

The monoclonal antibody of claim 11 (which is independent of claim 1), 12, 16 and 17 is not novel with regard to D3 (Art. 33 (2) PCT).

Moreover, the subject matter claims 20, 25, 26, 28, 29, 31-35, 38, 43-46 and 49 lacks novelty with regard to the disclosures in D1 or D2 or D3 (Art. 33 (2) PCT).

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**INTERNATIONAL PRELIMINARY  
EXAMINATION REPORT - SEPARATE SHEET**

---

International application No. PCT/GB97/01161

- 4). The problem underlying the present application, namely the appearance of escape mutants and the detection and prevention of HBV escape mutant infections, is already recognized in the prior art (D1/D2, D3) which also offers solutions. Therefore the subject matter of claims 9, 18, 19, 24, 27, 30, 36, 37, 39-42, 47, 48, 50-58 is at least obviously derivable from D1/D2 or D3 and cannot be called inventive (Art. 33 (3) PCT).
- 5). The specific deposited monoclonal antibodies and the hybridomas producing them ( claims 13-15 and 21-23) appear to be novel and not obviously derivable from the prior art cited in the International Search Report (Art. 33 (2), Art. 33 (3) PCT).
- 6). For the assessment of the present claims 40, 52 and 58 on the question whether they are industrially applicable, no unified criteria exist in the PCT. The patentability can also be dependent upon the formulation of the claims. The EPO, for example, does not recognize as industrially applicable the subject-matter of claims to the use of a compound in medical treatment, but may allow, however, claims to a known compound for first use in medical treatment and the use of such a compound for the manufacture of a medicament for a new medical treatment.

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